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____ Examiner # :_____ Date: _____

Serial Number:

SEARCH REQUEST FORM

Scientific and Technical Information Center

Art Unit: Pho	one Number 30	Serial Number:
Mail Box and Bldg/Room Loc	ation:	Results Format Preferred (circle): PAPER DISK E-MAI
If more than one search is s	ubmitted, please p	rioritize searches in order of need. ***********************************
Please provide a detailed statement	of the search topic, and dures, keywords, synonym terms that may have a sp	escribe as specifically as possible the subject matter to be searched. is, acronyms, and registry numbers, and combine with the concept or ecial meaning. Give examples or relevant citations, authors, etc, if
Inventors (please provide full nam		
Earliest Priority Filing Date:		
For Sequence Searches Only Pleas appropriote serial number.	e include all pertinent info	rmation (porent, child, divisional, or issued patent numbers) along with the
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	**************************************	ch Vendors and cost where applicable
STAFF USE ONLY Searcher: 0. Schrebe		•••
Searcher Phone #: 308 - 42		
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Date Searcher Picked Up:	Bibliographic	
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Searcher Prep & Review Time:		Sequence Systems Country 989
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Online Time:	Other	Other (specify)

Requester's Full Name: _

STIC-Biotech/ChemLib

From: Sent: To:

Seharaseyon, Jegatheesan Wednesday, May 02, 2001 3:59 PM STIC-Biotech/ChemLib Re:09/467160

Subject:

Please search the commercial database for SEQ ID:3 of 09/467,160. Thanks.

J.Seharaseyon Art Unit 1647 CM1 10D16 10CO1 MB (703)-305-1112

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(without alignments)
224.713 Million cell updates/sec
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379
1 APLATELRCQCLQTLQGIHL.....PASPMVKKIIEKMLKNGKSN
                                                                                                                May 4, 2001, 07:35:28; Search time 18.57 Seconds
GenCore version 4.5
Copyright (c) 1993 - 2000 Compugen Ltd.
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Total number of hits satisfying chosen parameters:

Minimum DB seq length: 0 Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

A_Geneseq_0401:* Database :

/SIDS6/gcgdata/geneseq/geneseqp/AA1982.DAT:*/SIDS6/gcgdata/geneseq/geneseqp/AA1983.DAT:*/SIDS6/gcgdata/geneseq/geneseqp/AA1984.DAT:*/SIDS6/gcgdata/geneseq/geneseqp/AA1985.DAT:*/SIDS6/gcgdata/geneseq/geneseqp/AA1985.DAT:*/SIDS6/gcgdata/geneseq/geneseqp/AA1985.DAT:*/SIDS6/gcgdata/geneseq/geneseqp/AA1987.DAT:*/SIDS6/gcgdata/geneseq/geneseqp/AA1988.DAT:*/SIDS6/gcgdata/geneseq/geneseqp/AA1988.DAT:* /SIDS6/gcgdata/geneseq/geneseqp/AA1980.DAT:* /SIDS6/gcgdata/geneseq/geneseqp/AA1981.DAT:* /SIDS6/gcgdata/geneseg/genesegp/AA1989.DAT /SIDS6/gcgdata/geneseg/genesegp/AA1991.DAT /SIDS6/gcgdata/geneseq/geneseqp/AA1999. /SIDS6/gcgdata/geneseq/geneseqp/AA1990. /SIDS6/gcgdata/geneseq/geneseqp/AA1995. /SIDS6/gcgdata/geneseq/geneseqp/AA1998. /SIDS6/gcgdata/geneseq/geneseqp/AA1994 /SIDS6/gcgdata/geneseq/geneseqp/AA1996 /SIDS6/gcgdata/geneseq/geneseqp/AA1997 /SIDS6/gcgdata/geneseq/geneseqp/AA1992 /SIDS6/gcgdata/geneseq/geneseqp/AA1993

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

/SIDS6/gcgdata/geneseq/geneseqp/AA2000.DAT: /SIDS6/gcgdata/geneseq/geneseqp/AA2001.DAT:

SUMMARIES

Description	MIP-2alpha. Homo	Human gro-beta che	Human chemokine gr	Human chemokine gr	Human gro-beta pro	human gro-beta pol	Amino acid sequenc	Human macrophage i	Human macrophage i	Human Gro beta cyt	Growth related onc
ID		R66699	W18025	W17671	X18104	W81499	X69036	R20589	R20529	R23034	W96714
DB	14	16	18	18	20	20	21	13	13	13	20
% Query Match Length DB	73	73	73	73	73	73	73	107	107	107	107
% Query Match	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0
Score	379	379	379	379	379	379	379	379	379	379	379
Result No.		2	m	4	Ŋ	9	7	ω	σ	10	11

Human chemokine GR Gro-beta/MIP-2-alp Human gro-alpha ch Human chemokine gr Human chemokine gr Human aro-alpha pr		Human chemokine GR Melanoma growth st MIP-2beta. Homo s Human gro-gamma ch	Protein used to ge Human chemokine gr Human gro-gamma pr Human mature gro-q	Amino acid sequenc Human Gro gamma cy Growth related onc Human macrophage i Gro-gamma/MIP-2-be	Human macrophage i Human macrophage i Cytokine and neutr Chimeric interleuk Mouse KC chemokine	Murine chemokine K Murine chemokine K Murine protein KC. Mouse mature KC po Chimeric interleuk Macrophage derived
B15810 R70793 R66698 W18024 W17670	W81498 Y69026 W96713 B15793	B15813 R70792 R36772 R66700	R93194 W18026 Y18107 W81500	Y69037 R23035 W96715 R20590 R70794	W1/0/2 R20530 R14077 W12434 R66697	W18023 W17669 Y18105 W81497 W12436 R05790
21 16 16 18 18	20 20 21 20 20 20 20 20 20 20 20 20 20 20 20 20	21 16 14	17 18 20 20	20 13 13 16	13 13 18 19	18 20 20 11 11
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379 331 331 331	331 331 331 331	331 330 325 325	325 325 325 325	9 9 9 9 9 9 9 9 9 9 9 9 9 9 9 9 9 9 9	3.23 3.18 2.84 2.83 2.78	278 278 278 278 267.5
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ALIGNMENTS

RESULT R36771	RESULT 1 - R36771	
QI X	R36771 standard; protein; 73 AA.	
XX	R36771;	
E X	29-SEP-1993 (first entry)	
XE	MIP-2alpha.	
ΚĚ		yocytopoiesis; MIP-1; MIP-2;
K K		; stroke; emboli; platelet;
XX		
SOX	Homo sapiens.	
Z Z	WO9309799-A.	
۲a:	27-MAY-1993.	
PF YX	13-NOV-1992; 92WO-US09671.	
Y K	15-NOV-1991; 91US-0792988.	
X & X	(UYPE-) UNIV PENNSYLVANIA.	
E.	Gewirtz AM;	
X Z X	WPI; 1993-182239/22.	
£ £ ;	Suppression of megakaryocytopoiesis - macrophage inflammatory protein-1 or -	by administration of 2
PS	Disclosure; Page 18; 26pp; English.	

73 AA;

Sequence

X S

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A claimed method for reducing the no. of circulating platelets in the bloodstream of a mammal comprises admin. of MIP-1 and/or MIP-2 or their analogues to induce such a reduction. The MIP-1, MIP-2 or analogue may be operatively linked to a carrier. The MIPs can be used to inhibit megakaryocytopoiesis to effect in vivo reduction of platelet nos. They can be used to treat disorders with excessively high platelet counts such as thrombocytosis, stroke, pulmonary emboli and myeloproliferative disorders.
*55555555555
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73 AA; Sequence

Gaps ; 0 Length 73; 100.0%; Score 379; DB 14; 100.0%; Pred. No. 1.4e-38; iive 0; Mismatches 0; Query Match 100. Best Local Similarity 100. Matches 73; Conservative

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1 APLATELRCQCLQTLQGIHLKNIQSVKVKSPGPHCAQTEVIATLKNGQKACLNPASPMVK 60

61 KIIEKMLKNGKSN 73

g

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RESULT R66699

R66699 standard; protein; 73 AA. R66699

19-JUL-1995 (first entry) Human gro-beta chemokine

hematopoietic synergistic factor; HSF; hematopoietic precursor; bone marrow; intercrine; desamino gro-beta; truncation. Gro-alpha protein; chemokine; inflammation; antiinflammatory;

Homo sapiens

Location/Qualifiers Protein

/note= "desamino truncated gro-beta, claim 6,

WO9429341-A

22-DEC-1994

94WO-US06264. 03-JUN-1994; 93US-0073800 08-JUN-1993; (SMIK) SMITHKLINE BEECHAM CORP.

Pelus LM; Bhatnagar P, King AG, WPI; 1995-036402/05. Balcarek JM,

truncated chemokine with increased biological activity - and related multimers, nucleic acid, antibodies etc., for treating inflammation, stimulating growth of bone marrow etc., also peptide(s) for inducing haematopoietic synergistic factor.

Disclosure; Page 51-52; 89pp; English

human mature gro-beta (full sequence given in R66699) shows enhanced biological activity when compared to the mature protein, and has been used to prepare multimeric, modified chemokines.

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A claimed method of treating or preventing sepsis comprises administering to an animal an effective amount of a chemokine selected from mature murine KC (see W18023), human gro plpha (see W18024), human gro beta (W18025) and human gro gamma (see W18026), modified fragments of these chemokines and multimeric proteins comprising an association of two chemokine proteins.

Sepsis can occur in hospitalised patients, and a consequence of bacterial sepsis is septic shock. The method of the invention provides a treatment for sepsis, particularly septic shock, which is major cause of death in intensive care units. Septic shock is yndrome apparently has intractable resistance to the effects of a variety of highly potent antimicrobial agents. Survival is increased by treatment with the chemokines, both prophylactically
                                                                       1 APLATELRCQCLQTLQGIHLKNIQSVKVKSPGPHCAQTEVIATLKNGQKACLNPASPMVK 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Use of chemokine(s) such as KC and gro-alpha - to treat or prevent sepsis, particularly septic shock
                              ö
 Length 73;
                                                                                                                                                                                                                                                                                                                                            Sepsis; septic shock; therapy; gro beta; chemokine; human.
                                                                                                                                                                                                                                                                                                                                                                                                                                   KC
                                Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                  /note= "preferred modified fragment of
  (Claim 2)"
Score 379; DB 16;
Pred. No. 1.4e-38;
Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (SMIK ) SMITHKLINE BEECHAM CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 1; Page 19; 28pp; English.
100.0%; Sc
100.0%; Pr
tive 0;
                                                                                                                                                                                                                          standard; protein; 73 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     95US-0007425
 Query Match 100.
Best Local Similarity 100.
Matches 73; Conservative
                                                                                                                                                                                                                                                                                                               Human chemokine gro beta.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DeMarsh PL, Johanson KO;
                                                                                                                     61 KIIEKMLKNGKSN 73
                                                                                                                                      WPI; 1997-298111/27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          and after infection
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                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                         Key
Protein
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TD W18025
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us-09-467-160-3.rag

agents

Gaps

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Indels

Length 73;

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This polypeptide sequence comprises human gro beta. Use of mammalian chemokines selected from gro alpha (W17670), gro beta, gro gamma (W17672) or KC (W17669) for mobilising haematopoietic cells is claimed. The chemokines, optionally used together with a growth factor or other haematopoietic regulator, are used to treat myelosuppression or any condition that involves immunosuppression or any condition haematopoietic disorders, aplastic anaemia or autoimmune disease, or low production/differentiation of haematopoietic or bone marrow cells. A claimed application is in peripheral blood stem cell transplants in patients being treated by chemotherapy. Chemokines having an N-terminal deletion are more active, e.g. by 2 orders of magnitude, than the full-length protein.

When used with a colony stimulating factor (CSF), a synergistic effect is achieved and the dose, and side-effects, of CSF can be reduced. Compared with conventional methods of stem cell transplants, reduced.
                                                                       1 APLATELRCQCLQTLQGIHLKNIQSVKVKSPGPHCAQTEVIATLKNGQKACLNPASPMVK 60
                                                                                            immunosuppression; haematopoietic cell; infection; cancer; aplastic anaemia; autoimmune disease; stem cell transplant;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /note= "preferred polypeptide (Claim 2)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Use of specific chemokine(s) for stem cell mobilisation treating myelosuppression or any condition involving immunosuppression or low levels of haematopoietic cells
                                                                                                                                                                                                                                                                                                                                                                                                                         Gro beta; chemokine; intercrine; myelosuppression;
100.0%; Score 379; DB 18;
100.0%; Pred. No. 1.4e-38;
ive 0; Mismatches 0;
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                                                                                                                                                                                                                                                                          W17671 standard; Protein; 73 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 1; Page 21; 31pp; English.
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                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                       Human chemokine gro beta.
                                     Conservative
                                                                                                                                                                 61 KIIEKMLKNGKSN 73
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 1997-258957/23.
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                  Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    24-OCT-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens
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                    Best Local Simi
Matches 73;
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 Query Match
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                                                                                                                                                                                                                                                                                                                                                             Human; gro-beta; haematopoietic stem cell mobilisation; myelosuppression; stem cell transplantation; haematopoietic recovery; therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          This sequence represents the human gro-beta protein. The invention relates to a method for mobilising haematopoietic stem cells comprising administering a modified gro-beta protein. The new method is useful for the treatment of patients receiving a peripheral blood haematopoietic stem cell transplantation, and for treatment of myelosuppression. The new method enhances haematopoietic recovery, as this still causes morbidity and mortality.
                                                                                      Gaps
                                                                                                                1 APLATELRCQCLQTLQGIHLKNIQSVKVKSPGPHCAQTEVIATLKNGQKACLNPASPMVK 60
                                                                                                                             1 APLATELRCQCLQTLQGIHLKNIQSVKVKSPGPHCAQTEVIATLKNGQKACLNPASPMVK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Use of modified chemokine to mobilize haematopoietic stem cells,
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                                                             Length 73;
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                                                            100.0%; Score 379; DB 18;
100.0%; Pred. No. 1.4e-38;
ive 0; Mismatches 0;
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                                                                                                                                                                                                                                                          Y18104 standard; protein; 73 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            in cell transplants
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                                                              Query Match 100.
Best Local Similarity 100.
Matches 73; Conservative
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Best Local Similarity 100.
Matches 73; Conservative
                                                                                                                                                                                                                                                                                                                                      Human gro-beta protein.
                                                                                                                                                                                61 KIIEKMLKNGKSN 73
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                        73 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             King AG, Qian Y;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  73 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                              WO9926645-Al.
                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens.
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useful for

use of mammaolian chemokines provides a more rapid release of peripheral blood mononuclear cells and/or platelets, reducing risk of infection and allowing use of higher doses of chemotherapeutic

61 kiiekmlkngksn 73

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This is the amino acid sequence of the human chemokine gro-beta mature polypeptide. The invention relates to a method of preventing and treating sepsis using chemokines selected from mature or modified murine KC (see WB1497); or human gro-alpha (see WB1498), gro-beta or gro-gamma (see WB1499). A claimed method uses a dimeric chemokine consisting of 2 covalently linked modified gro-beta proteins (amino acids 5-73 of the mature polypeptide) in which the proteins are linked by 2 intermolecular disulphide bonds between C5-C31 and C7-C47. Further claimed is administering the chemokine in conjunction with an anti-infective agent. The chemokines described in this invention are required to treat and abrogate septic mortality.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 APLATELRCQCLQTLQGIHLKNIQSVKVKSPGPHCAQTEVIATLKNGQKACLNPASPMVK 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Treating and preventing sepsis in animals – by using two modified gro b chemokines in conjunction with an anti-infective agent
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 73;
                                                                                                                                                Gro-beta; chemokine; human; sepsis; septic shock; therapy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               100.0%; Score 379; DB 20;
100.0%; Pred. No. 1.4e-38;
ive 0; Mismatches 0;
                                                                                                                     human gro-beta polypeptide used to treat sepsis.
                                                                                                                                                                                                                          5..73
/note= "claimed fragment"
                                                                                                                                                                                                              Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 1; Page 18; 26pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Y69036
ID Y69036 standard; protein; 73 AA.
                                                                                                                                                                                                                                                                                                                                                                                          (SMIK ) SMITHKLINE BEECHAM CORP.
                          W81499 standard; Protein; 73 AA
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                                                                                                                                                                                                                                                                                                                                                             97US-0846966
                                                                                      01-MAR-1999 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                         Demarsh PL, Johanson KO;
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Best Local Similarity
Las 73; Conserve
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                                                                                                                                                                                                                                                                                                                                                             29-APR-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence
                                                                                                                                                                                                                            Protein
                                                                                                                                                                                Mus sp.
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The present sequence represents a chemokine receptor ligand. The present ligand can be incorporated into the conjugates of the invention. The specification describes a conjugate, comprising a targeted agent and a chemokine receptor ligand. The conjugate binds to a chemokine receptor receptor ligand. The conjugate binds to a chemokine receptor receptor. The conjugates are used for formulating a medicament or for treating disorders associated with inflammatory responses resulting from activation, proliferation and migration of immune effector cells. The disorders or disease states comprise secondary tissue damage such as central nervous system (CNS) injury, CNS inflammatory diseases, neurodegenerative disorders, heart disease, inflammatory eye diseases, inflammatory bowel diseases, inflammatory line and the conjugate of the conju
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A new therapeutic agent comprising a conjugate for treating secondary tissue damage and other disease conditions like Alzheimer's disease, stroke, Parkinson's disease and atherosclerosis
                                                                                                                                                                                             Chemokine receptor; ligand; inflammatory response; immune effector ce secondary tissue damage; central nervous system injury; MTP-Zalpha; CNS inflammatory disease; neurodegenerative disorder; heart disease; inflammatory eye disease; inflammatory bowel disease; inflammatory joint disease; inflammatory kidney; renal disease; inflammatory lung disease; inflammatory masal disease; inflammatory huy disease; inflammatory huyoid disease; thyroiditis; cytokine regulated cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         or renal diseases, inflammatory lung diseases, inflammatory diseases, inflammatory thyroid disease such as thyroiditis, or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 APLATELRCQCLQTLQGIHLKNIQSVKVKSPGPHCAQTEVIATLKNGQKACLNPASPMVK 60
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                                                                                                                                          Amino acid sequence of chemokine receptor ligand MIP-2alpha.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Disclosure; Page 60; 204pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               99WO-CA00659.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           cytokine-regulated cancers.
                                                                                 30-MAY-2000 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Coggins PJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (OSPR-) OSPREY PHARM LTD.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2000-182542/16.
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                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens
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1 APLATELRCQCLQTLQGIHLKNIQSVKVKSPGPHCAQTEVIATLKNGQKACLNPASPMVK 60
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       genes can be used to produce recombinant MIP proteins for use in wound healing, to modulate myelopoiesis and to induce adjuvant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human macrophage inflammatory protein 2-alpha - useful in
treating infections, cancer, myelopoletic dysfunction and auto:
immune diseases
                                                     Inducible; secretory; inflammatory response; cytokine.
                 Human macrophage inflammatory protein (MIP) 2 alpha
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            activity.
See also R20588,R20590 and R20528-30.
                                                                                                                              Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure; Fig 2; 68pp; English.
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/label= signal
                                                                                                                                                                                                                                                                                                                                                                                                                               Tekamp-Olson P, Gallegos CA;
                                                                                                                                                                                                                                                                                                91WO-US04482.
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90US-0541897
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 1992-041519/05
                                                                                                                                                                                                                                                                                                                                                                                          (CHIR-) CHIRON CORP
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                                                                                         Homo sapiens
                                                                                                                                                                                                                                                                                                24-JUN-1991;
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                                                                                                                                                                                                                                                                                                                                                        22-JUN-1990;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        the U937 cDNA library prepd. from poly-A+ RNA from PMA treated and LPS stimulated cells, using as probe a fragment isolated from the mMIP-2 cDNA (see Q20612) encoding most of the mature mMIP-2 amino acid sequence. Two classes of human cDNA homolgous to mMIP-2 were
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The sequence was deduced from the DNA sequence obtd. by screening
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           genes can be used to produce recombinant MIP proteins for use in wound healing, to modulate myelopoiesis and to induce adjuvant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   found designated alpha and beta. The alpha form reproduced here is claimed in W09200327; the beta form (020614) is claimed in W09200326. The alpha form is the more abundant of the two. The genes can be used to produce recombinant MIP proteins for use in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human macrophage inflammatory protein 2-alpha - useful in treating infections, cancer, myelopoietic dysfunction and auto:
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                                                                                                           Inducible; secretory; inflammatory response; cytokine
                                                                    Human macrophage inflammatory protein (MIP) 2 alpha.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  See also R20588, R20590 and R20528-30.
                                                                                                                                                                                 Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; Fig 2; 68pp; English.
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/label= signal
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                                                                                                                                                                                                                                                                                                                                                      91WO-US04478
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                                23-APR-1992 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                (CHIR-) CHIRON CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 1992-041518/05
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Matches 73; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       107 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   immune diseases
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22-JUN-1990;
                                                                                                                                               Homo sapiens
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Gaps

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Query Match

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20 x 8

R20529;

RESULT R20529

Length 107; Indels

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The cDNA clone coding for inflammatory cytokine Gro-beta was isolated from a mezerin- and calcium ionophore-stimulated leukocyte cDNA library. The amino acid sequence of Gro beta was deduced from the nucleotide sequence. The level of Gro beta in inflammatory response cells can be used as an indication of a test substance's inflammatory activity and to diagnose colon cancer.
                                                                                                                                                                                                                                                                                                                               beta and Gro gamma inflammatory cytokine(s) - for use in
Cytokine; inflammatory response; MAD-2; cancer diagnosis; colonic epithelial tumour cell.
                                                         Location/Qualifiers
                                                                                         /note= "putative"
36..107
/label= Gro_beta
                                                                                                                                                                                                                                                                                                                                                                  Claim 12; Fig 1B; 46pp; English.
                                                                                                                                                                                                                                   (CETU ) CETUS CORP.
(UYNC-) UNIV OF NORTH CAROLINA.
                                                                     1..35
/label= leader
                                                                                                                                                                                       91WO-US06936.
                                                                                                                                                                                                             90US-0590223
                                                                                                                                                                                                                                                                      Haskill JS, Nitecki DE,
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N-PSDB; Q24266.
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                                                                                                                                                                                                                                                                                                                                             diagnosing colon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                See also Q24267
                                                                                                                                                                                       24-SEP-1991;
                                   Homo sapiens
                                                                                                                                                                                                             28-SEP-1990;
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                                                                                                                                                               16-APR-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence
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Ralph P;

ö Gaps ö Length 107; Indels 100.0%; Score 379; DB 13; 100.0%; Pred. No. 2.3e-38; iive 0; Mismatches 0; 73; Conservative

1 APLATELRCQCLQTLQGIHLKNIQSVKVKSPGPHCAQTEVIATLKNGQKACLNPASPMVK 60

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W96714 standard; Protein; 107 AA. 15-APR-1999 (first entry) W96714;

Growth related oncogene-beta; GRO-beta; CXC chemokine; metastasis;

Growth related oncogene-beta (GRO-beta).

anglogenesis inhibitor; angiostasis inducer; tumour growth inhibition; haemangiomas; rheumatoid arthritis; atherosclerosis; meningioma; idopathic pulmonary fibrosis; benjon prostatic hypertrophy; psoriasis; vascular restenosis; arteriovenous malformation; neovascular glaucoma; anglofibroma; haemophilic joint; hypertrophic scar; Osler-Weber syndrome;

The present sequence is a growth related oncogene-beta (GRO-beta). GRO-alpha is a CXC chemokine that is an inhibitor of angiogenesis. The specification describes methods for inhibiting angiogenesis or for inducing angiostasis, using chemokines (with a conserved Cys Xaa Cys (CXC) sequence at the N-terminal) other than platelet factor-4, and which do not contain the amino acid sequence EIR. The methods are useful for inhibiting tumour growth and metastasis and for treating diseases such as haemangiomas, rheumatoid arthritis, atherosclerosis and idiopathic pulmonary fibrosis (IPF), benign prostatic hypertrophy (BPH), vascular restenosis, arteriovenous malformations (AVM), meningioma, meovascular glaucoma, psoriasis, angiofibroma, haemophilic johits, hypertrophic scars, Osler Weber syndrome, pyogenic granuloma retrolental fibroplasia, scleroderma, trachoma, vascular adhesions, synovitis, febroplasis, pterygium, diabetic retinopathy neovascularisation associated with corneal injury or grafts, adult respiratory distress syndrome (ARDS), chronic bronchitis, pseudogout pterygium; Use of chemokines with a conserved Cys Xaa Cys (CXC) sequence which do not contain amino acid sequence ELR, for inhibiting angiogenesis in tumours, rheumatoid arthritis, restenosis or pyogenic granuloma retrolental fibroplasia; scleroderma; trac. vascular adhesion; synovitis; dermatitis; endometriosis; pterdiabetic retinopathy; neovascularisation; chronic bronchitis; adult respiratory distress syndrome; ARDS; pseudogout; Disclosure; Columns 95-96; 145pp; English. RM; Strieter 95US-0468819 Polverini PJ, (UNMI) UNIV MICHIGAN and cystic fibrosis. WPI; 1999-166569/14. 107 AA; cystic fibrosis N-PSDB; X15002 Homo sapiens 06-JUN-1995; US5871723-A. 16-FEB-1999 Kunkel SL, Sequence glaucoma MAKE A MA

Gaps ö Length 107; Indels ö Score 379; DB 20; Pred. No. 2.3e-38; Mismatches 100.0%; Sc 100.0%; Pr tive 0; Conservative Local Similarity 73; Query Match Best Loca Matches

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1 APLATELRCQCLQTLQGIHLKNIQSVKVKSPGPHCAQTEVIATLKNGQKACLNPASPMVK 60

> ò g

61 KIIEKMLKNGKSN 73 ò

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B15810 standard; Protein; 107 B15810; B15810

RESULT 12

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17-JAN-2001 (first entry)

Human chemokine GRObeta SEQ ID NO: 53.

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                                          monocyte chemoattractant protein-1; inflammation; atherosclerosis; HIV; HIDS; stroke; psoriasis; autoimmune disease; hypertension; endotoxaemia; basophil-mediated disease; myocardial infarction; acute ischaemia; rheumatoid arthritis; contraception.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           chemokines which can be used to produce derivatives, agonists and antagonists which are then useful in disease treatment. The chemokines include sequences B15785-B15794, B15803-B15813 and B15813-B15840. These chemokine derivatives can be used to treat diseases such as autoimmune diseases, atherosclerosis, osteoporosis, HIV infection and AIDS, psoriasis, inflammatory diseases, hypertension, basophil-mediated diseases, endotroxaemia, myocardial infarction, acute ischaemia and rheumatoid arthritis, and can be used to prevent strokes and as contraceptives. The coding sequences for the chemokines can be used in gene therapy for the same diseases, as well as in the production of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New peptide 3, amide and heterocyclic compounds and saccharide conjugates used for inhibiting chemokine induced activity and for treating e.g. stroke, vascular diseases, autoimmune diseases and tumour
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 APLATELRCOCLQTLQGIHLKNIQSVKVKSPGPHCAQTEVIATLKNGQKACLNPASPMVK 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Macrophage recruitment; chemokine derivative; MCP-1; osteoporosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present invention concerns the identification of a number of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ő
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Macrophage inflammatory protein 2-alpha; gro-beta/MIP-2-alpha;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      100.0%; Score 379; DB 21; Length 107; 100.0%; Pred. No. 2.3e-38; 0; Mismatches 0; Indels 0.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Disclosure; Page 364; 387pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  R70793 standard; Protein; 102 AA.
                                                                                                                                                                                                                                                                                                                                         99US-0229071.
99US-0271192.
99US-0452406.
                                                                                                                                                                                                                                                                                               12-JAN-2000; 2000WO-US00821
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Grainger DJ, Tatalick LM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match 100.
Best Local Similarity 100.
Matches 73; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    kiiekmlkngksn 107
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          KIIEKMLKNGKSN 73
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gro-beta/MIP-2-alpha.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2000-499101/44.
N-PSDB; A74871.
                                                                                                                                                                                                                                                                                                                                                                                                                                 (NEOR-) NEORX CORP.
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                                                                                                                                                                                                      WO200042071-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             animal models
                                                                                                                                                          Homo sapiens
                                                                                                                                                                                                                                                                                                                                           12-JAN-1999;
17-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                       01-DEC-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             29-AUG-1995
                                                                                                                                                                                                                                                  20-JUL-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                R70793;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  growth
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               R70793
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Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                Screening for cpds. with anti-heparanase activity - by detecting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            hematopoietic synergistic factor; HSF; hematopoietic precursor;
bone marrow; intercrine; desamino gro-alpha; truncation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            5..732
/note= "desamino truncated gro-alpha, claim 7,
page 69"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                inhibition of heparin or heparan sulphate degradation, potentially useful for treating arthritis, restenosis, cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gro-alpha protein; chemokine; inflammation; antiinflammatory;
heparanase; heparin; heparan sulfate; arthritis; restenosis; cancer; wound healing.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 102;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Pred. No. 3.9e-35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 12; Page 40; 60pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  R66698 standard; protein; 73 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            100.08;
                                                                                                                                                                                                                                         93US-0099866
93US-0136117
                                                                                                                                                                                              94WO-US08207
                                                                                                                                                                                                                                                                                                                                                       SR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human gro-alpha chemokine..
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     19-JUL-1995 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Similarity 100.
68; Conservative
                                                                                                                                                                                                                                                                                                                                                       Ledbetter
                                                                                                                                                                                                                                                                                                                                                                                               WPI; 1995-082239/11.
                                                                                                                                                                                                                                                                                                            (UPJO ) UPJOHN CO.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    102 AA;
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                                                                   Homo sapiens
                                                                                                                                                                                              26-JUL-1994;
                                                                                                                                                                                                                                         29-JUL-1993;
13-OCT-1993;
                                                                                                                                                                                                                                                                                                                                                       Hoogwerf AJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens
                                                                                                         WO9504158-A
                                                                                                                                                    09-FEB-1995
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Matches 6
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Protein
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R66698
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Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          Truncated, desamino chemokine comprising amino acids 5-73 of human mature gro-alpha (full sequence given in R66698) shows enhanced biological activity when compared to the mature protein, and has been used to prepare multimeric, modified
                                                                                                                                                              New truncated chemokine with increased biological activity - ar related multimers, nucleic acid, antibodies etc., for treating inflammation, stimulating growth of bone marrow etc., also
                                                                                                                                                                                                                                                                                                                                                                                     Query Match 87.3%; Score 331; DB 16; Length 73; Best Local Similarity 87.7%; Pred. No. 8.8e-33; Matches 64; Conservative 4; Mismatches 5; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sepsis; septic shock; therapy; gro alpha; chemokine; human.
                                                                                                                                                                                                         peptide(s) for inducing haematopoietic synergistic factor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /note= "preferred modified fragment of KC
                                                                                                            Pelus LM;
                                                                                                                                                                                                                                                                                                                                                                                                  Pred. No. 8.86
4; Mismatches
                                                                                                         Bhatnagar P, King AG,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Location/Qualifiers
                                                                                                                                                                                                                                  Disclosure; Page 51; 89pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (Claim 4)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (SMIK ) SMITHKLINE BEECHAM CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               W18024 standard; protein; 73 AA.
                                                                               (SMIK ) SMITHKLINE BEECHAM CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          96WO-US18616
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                           94WO-US06264
                                                    93US-0073800
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61 kiiekmlnsdksn 73
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                                                                                                                                      WPI; 1995-036402/05
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                                                                                                                                                                                                                                                                                                                                               73 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens
                         03-JUN-1994;
                                                    08-JUN-1993;
                                                                                                           Balcarek JM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO9719173-A1
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                                                                                                                                                                                                                                                                                                                     chemokines
                                                                                                                                                                                                                                                                                                                                               Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          W18024;
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W18024
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A claimed method of treating or preventing sepsis comprises administering to an animal an effective amount of a chemokine selected from mature murine KC (see W18023), human gro alpha (W18024), human gro beta (see W18025) and human gro gamma (see W18025), modified fragments of these chemokines and multimeric proteins comprising an association of two chemokine proteins. Sepsis can occur in hospitalised patients, and a consequence of bacterial sepsis is septic shock. The method of the invention provides a treatment for sepsis, particularly septic shock, which is a major cause of death in intensive care units. Septic shock is syndrome apparently has intractable resistance to the effects of a variety of highly potent antimicrobial agents. Survival is increased by treatment with the chemokines, both prophylactically
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 APLATELRCQCLQTLQGIHLKNIQSVKVKSPGPHCAQTEVIATLKNGQKACLNPASPMVK 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   of chemokine(s) such as KC and gro-alpha - to treat or prevent is, particularly septic shock
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 73;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 331; DB 18;
Pred. No. 8.8e-33;
4; Mismatches 5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Search completed: May 4, 2001, 07:37:02 Job time: 94 sec
                                                                    Claim 1; Page 18-19; 28pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                    87.3%;
87.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         64; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        61 KIIEKMLKNGKSN 73
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kiiekmlnsdksn 73
                                                                                                                                                                                                                                                                                                                                                          and after infection
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                             73 AA;
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                                     sepsis,
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Query Match
Best Local Similarity
Matches 73; Conserv
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                                                                                                                                                                                                                                                                                                                                         RESULT
                                                                                       May 4, 2001, 07:35:28; Search time 11.75 Seconds (without alignments) 119.353 Million cell updates/sec
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Sequence 3, M
Sequence 4, M
Sequence 6, M
Sequence 4, M
Sequence 4, M
Sequence 3, M
Sequence 3, M
Sequence 3, M
Sequence 18, M
Sequence 18, M
Sequence 18, M
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Sequence 27,
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                                                                                                                                                                                 ........PASPMVKKIIEKMLKNGKSN
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/cgn2_6/ptodata/2/jaa/5B_COMB.pep:*
/cgn2_6/ptodata/2/jaa/6A_COMB.pep:*
/cgn2_6/ptodata/2/jaa/pcUUS_COMB.pep:*
/cgn2_6/ptodata/2/jaa/pcTUS_COMB.pep:*
/cgn2_6/ptodata/2/jaa/pcTUS_COMB.pep:*
4.5
Compugen Ltd
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US - 08 - 352 - 324A - 4
US - 08 - 862 - 607 - 4
US - 08 - 468 - 819 - 6
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US-08-482-111-3
US-07-778-413E-18
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US-08-330-163-2
US-08-482-111-2
US-08-436-420-26
                                                                                                                                                                                                                                                                                          Total number of hits satisfying chosen parameters:
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                                                                                                                                                                                                                                                            185757 segs, 19210857 residues
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GenCore version
Copyright (c) 1993 - 2000
                                                                                                                                                                                                                                                                                                                                                                                                    Listing first 45 summaries
                                                          protein search, using sw model
                                                                                                                                                                                                                 BLOSUM62
Gapop 10.0 , Gapext 0.5
                                                                                                                                                                   379
1 APLATELRCQCLQTLQGIHL.
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                                                                                                                                                                                                                                                                                                                                                                      Post-processing: Minimum Match 0% Maximum Match 100%
                                                                                                                                                                                                                                                                                                                          Minimum DB seq length: 0
Maximum DB seq length: 2000000000
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Match Length
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100.0
100.0
96.8
96.8
87.3
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                                                            OM protein
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                                                                                                                                                                                   Sequence:
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                                                                                          Run on:
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Appli
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APPLICANT: Gewirtz, Alan M.
TITLE OF INVENTION: Suppression of megakaryo-
TITLE OF INVENTION: cytopoiesis by macrophage inflammatory proteins
Sequence 20, Sequence 20, Sequence 20, Sequence 4, P. Sequence 4, P. Sequence 4, P. Sequence 5, A. Sequence 1, 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 73;
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llarity 100.0%; Pred. No. 1.7e-40;
Conservative 0; Mismatches 0;
US-07-792-988-4
US-08-340-102-20
US-08-330-163-4
US-08-482-111-4
US-08-482-111-4
US-08-482-111-4
US-08-482-111-4
US-08-482-111-4
US-08-846-966-4
US-08-95-142-4
US-08-95-142-4
US-08-95-142-4
US-08-95-16144-5
US-08-48-19-7
US-09-203-235-5
US-08-48-19-7
US-09-203-235-5
US-08-48-19-7
US-09-203-235-5
US-08-48-19-7
US-08-48-19-7
US-09-203-235-5
US-08-48-40-38
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY: U.S.A.
ZIP: 19104-3246
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.50 inch, 720
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     University of Pennsylvania
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MEDIUM 115...
COMPUTER: IBM PS/2
OPERATING SYSTEM: MS-DOS
SOFTWARE: WORDPERFECT 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/792,988
FILING DATE: 1991115
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADDRESSEE: University of Penns STREET: Suite 419
STREET: 133 South 36th Street CITY: Philadelphia STATE: Pennsylvania
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILING DATE: 19911115
CLASSIFICATION: 514
PRIOR APPLICATION 514
PRIOR APPLICATION DATA:
APPRICATION NUMBER:
FILING DATE:
ATTORNEY/AGENY INFORMATION:
NAME: MONACO, Daniel A.
REGISTRATION NUMBER: 30,480
REFERENCE/DOCKET NUMBER: 6056
TELECOMMUNICATION INFORMATION:
TELECHONE: (215) 568-539
TELERAX: (215) 568-549
TELEX: NO. 5306709
INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: AMINO ACID
STRANDEDNESS: single stranded
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Patent No. 5306709
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     NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS
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PRIOR APPLICATION DATA
 CORRESPONDENCE ADDRESS:
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1 APLATELRCQCLQTLQGIHLKNIQSVKVKSPGPHCAQTEVIATLKNGQKACLNPASPMVK 60
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100.0%; Pred. No. 1.7e-40;
Live 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: No. 5591718el Neutrophil TITLE OF INVENTION: Activating Factors NUMBER OF SEQUENCES: 22
                                                                                                                                                                                                                                      APPLICANT: Walz, Alfred
TITLE OF INVENTION: No. 5401651e1 Neutrophil
TITLE OF INVENTION: Activating Factors
NUMBER OF SEQUENCES: 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ATTORNEY AGENT INFORMATION:
NAME: Cook, Robert R.
REGISTRATION NUMBER: 31602
REFERENCE/DCOKET NUMBER: A-204
TELECOMMUNICATION INFORMATION:
TELEPHONE: (865) 499-8011
INFORMATION FOR SEQ ID NO: 19:
                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.5 in., DS, 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER: Apple Macintosh
OPERATING SYSTEM: Macintosh OS 7.0
SOFTWARE: Microsoft Word Version 5.1a
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/778,413E
FILING DATE: 16-OCT-1991
CLASSIFICATION: 536
                                                                                                                                                                                  Sequence 19, Application US/07778413E Patent No. 5401651 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                        ADDRESSEE: Amgen Inc.
STREET: Amgen Center
STREET: 1840 Dehavilland Drive
CITY: Thousand Oaks
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 19, Application US/08340102
Patent No. 5591718
GENERAL INFORMATION:
APPLICANT: Walz, Alfred
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
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                                                                    61 KIIEKMLKNGKSN 73
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Best Local Similarity
Loca 73; Conserve
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    amino acid
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US-07-778-413E-19
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1 APLATELRCQCLQTLQGIHLKNIQSVKVKSPGPHCAQTEVIATLKNGQKACLNPASPWVK 60
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25 (EPO)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-08-436-420-27
Sequence 27, Application US/08436420
September 10 Sequence 30 and Applicant PROOST, Paul
TITLE OF INVENTION: GRANULOCYTE CHEMOTACTIC PROTEIN NUMBER OF SEQUENCES: 45
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       core 379; DB 1;
Pred. No. 1.7e-40;
; Mismatches 0:
                   STREET: 1840 Dehavilland Drive
CTYY: Thousand Oaks
STATE: California
COUNTRY: USA
ZIP: 91320-1789
COMPUTER REABABLE FORM:
MEDIUM TYPE: Diskette, 3.5 in., DS, 1.4 MB
COMPUTER: MS-DOS
                                                                                                                                                                                                            OPERATING SYSTEM: MS-DOS 6.22
SOFTWARE: Microsoft Word Version 5.1a for
SOFTWARE: Macintosh
                                                                                                                                                                                                                                                                                                                                                                                                                                             REFERENCE/DOCKET NUMBER: A-204A
TELECOMMUNICATION INFORMATION:
TELEPHONE: (805) 499-5725 EXTENSION 4955
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STREET: 1455 PENNSYLVANIA AVENUE, N.W. CITY: WASHINGTON STATE: DISTRICT OF COLUMBIA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              100.0%; Score 379;
                                                                                                                                                                                                                                                                                   CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/340,102
FILING DATE: 15-NOV-1994
CIASSIFICATION: 514
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US/08/436,420
FILING DATE: 24-MAY-1995
STREET: Amgen Center
STREET: 1840 Dehavilland Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADDRESSEE: HALE and DORR LLP
                                                                                                                                                                                                                                                                                                                                                                                                        NAME: Cook, Robert R. REGISTRATION NUMBER: 31602
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEFAX: (805) 499-8011
INFORMATION FOR SEQ ID NO: 19:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      100.08;
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MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 100.(
Best Local Similarity 100.(
Matches 73; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  73 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; MOLECULE TYPE: peptide US-08-340-102-19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                61 KIIEKMLKNGKSN 73
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                linear
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TYPE: amino acid
STRANDEDNESS:
TOPOLOGY: unknown
MOLECULE TYPE: protein
US-08-846-966-3
                      73 amino acids
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Best Local Similarity 100.
Matches 73; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEFAX: 215-270-5090 INFORMATION FOR SEQ ID NO:
      SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; MOLECULE TYPE: protein US-08-557-142-3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ZIP: 19406-2799
COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                       61 KIIEKMLKNGKSN 73
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILING DATE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-08-557-142-3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
ADDRESSE: SmithKline Beecham Corporation - Corporate Patents STREET: 709 Swedeland Road
CITY: King of Prussia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 73;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/846,966
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   100.0%; Score 379; DB 2;
100.0%; Pred. No. 1.7e-40;
tive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: SmithKline Beecham Corporation
APPLICANT: DeMarsh, Peter L.
APPLICANT: Johanson, Kyung O.
ATTLE OF INVENTION: Method of Treating Sepsis
NUMBER OF SEQUENCES: 4
                                           CLASSILLCALLON DATA:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/982,539
FILING DATE: 27-NOV-1992
ATTORNEY/AGENT INFORMATION:
NAME: BAKER HOLLIG L.
REGISTRATION NUMBER: 31,321
REFERENCE/DOCKET NUMBER: 102378.215
TELECHOME: (202) 942-8400
TELEFAX: (202) 942-8400
                                                                                                                                                                                  102378.215
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STAIL.

COUNTY: USA
ZIP: 1946-2799
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
"MADTER: IBM PC COMPATIBLE
"STAFFN: PC-DOS/MS-DOS
APPLICATION NUMBER: PCT/EP93/03330 FILING DATE: 26-NOV-1993 CLASSIFICATION: 530
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 60/007,425
FILING DATE: 21-NOV-1995
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         REFERENCE/DOCKET NUMBER: P50417-1
TELECOMMUNICATION INFORMATION:
TELEPHONE: 610-270-5016
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 3, Application US/08846966; Patent No. 6042821; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: Hall, Linda E. REGISTRATION NUMBER: 31,763
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            INFORMATION FOR SEQ ID NO: 3:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
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                                                                                                                                                                                                                                                                                                                                                                   TOPOLOGY: linear MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   61 KIIEKMLKNGKSN 73
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                                                                                                                                                                                                                                                                                                                              amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
Matches 73; Conserv
                                       CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                              STRANDEDNESS:
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                                                                                                                                                                                                                                                                                                                                                                                                           US-08-436-420-27
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US-08-846-966-3
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                                                                                              1 APLATELRCQCLQTLQGIHLKNIQSVKVKSPGPHCAQTEVIATLKNGQKACLNPASPMVK 60
                                                                                                                   Gaps
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  Length 73;
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                                                                                                                                                                                                                                                                                                                                            Sequence 3, Application US/08551142
| Patent No. 6080398
| GENERAL INFORMATION:
| APPLICANT: Pelus, Louis M
| APPLICANT: Bhatnagar, Pradip K
| APPLICANT: Bhatnagar, Pradip K
| APPLICANT: Blactarek, Joana M
| TITLE OF INVENTION: Methods of Enhancing Bioactivity of TITLE OF INVENTION: Chemokines
| NUMBER OF SEQUENCES: 29
| CORRESPONDENCE ADDRESS:
                                                 Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
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Query Match 100.0%; Score 379; DB 3; Best Local Similarity 100.0%; Pred. No. 1.7e-40; Matches 73; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   100.0%; Score 379; DB 3; 100.0%; Pred. No. 1.7e-40;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE: SmithKline Beecham Corporation
ADDRESSEE: Corporate Patents
STREET: 709 Swedeland Road
CITY: King of Prussia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ATTORNEY/AGENT INFORMATION:
NAME: Hall, Linda E.
REGISTRATION NUMBER: 31,763
REFERENCE/DOCKET NUMBER: SECP50161
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/073,800
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COMPUTER: IBM PC compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELECOMMUNICATION INFORMATION TELEPHONE: 215-270-5015
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ATTORNEY/AGENT INFORMATION:
NAME: Luther, Barbara J.
REGISTRATION NUMBER: 33,954
REFERENCE/DOCKET NUMBER: PF
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TELEPHONE: 415-855-0555
TELEFAX: 415-852-0195
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                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM COMPATIBLE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEFAX: 415-852-0195
INFORMATION FOR SEQ ID NO: 4:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   107 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity 100.0
Matches 73; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES: (CORRESPONDENCE ADDRESS)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                61 KIIEKMLKNGKSN 73
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       linear
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                                                                                                                                                                                                  94304
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                                                                                                                                                                  CA
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                                                                                                                                                                STATE: CA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 9
US-08-862-607-4
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                  1 APLATELRCQCLQTLQGIHLKNIQSVKVKSPGPHCAQTEVIATLKNGQKACLNPASPMVK 60
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                                                                                                                                                                                                                                APPLICANT: Pelus, Louis M
APPLICANT: Bhatnagar, Pradip K
APPLICANT: Ring, Andrew G
APPLICANT: Balcarek, Joanna M
TITLE OF INVENTION: Methods of Enhancing Bioactivity of
                                                                                                                                                                                                                                                                                                                                                                                  Corporate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ZIP: 194062799
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PCDOS/MSDOS
SOFWMARE: PATEAITIN Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US94/06264
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 100.0%; Score 379; DB 5; Best Local Similarity 100.0%; Pred. No. 1.7e-40; Matches 73; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES: 29
CORRESPONDENCE ADDRESS:
ADDRESSEE: SmithKline Beecham Corporation
ADDRESSEE: Patents
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ATTORNEY/AGENT INFORMATION:
NAME: Hall, Linda E.
REGIETRATION NUMBER: 31,763
REFERICE/DOCKET NUMBER: SBC P50161
TELECOMMUNICATION INFORMATION:
TELEPHONE: 215-270-5015
TELEFRAX: 215-270-5090
                                                                                                                                                                                                Sequence 3, Application PC/TUS9406264 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 4, Application US/08352324A; Patent No. 5633149; GENERAL INFORMATION:
APPLICANT: Guegler, Karl J.
                                                                                                                                                                                                                                                                                                                                                                                                                    709 Swedeland Road
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEFAX: 215-270-5090
INFORMATION FOR SEQ ID NO: 3:
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                                                                                                                                                                                                                                                                                                                                                                                                                                     King of Prussia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TOPOLOGY: unknown
MOLECULE TYPE: protein
                                                                                           61 KIIEKMLKNGKSN 73
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   61 KIIEKMLKNGKSN 73
                                                                     61 KIIEKMLKNGKSN 73
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                         PA
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US-08-352-324A-4
                                                                                                                                                             RESULT 7
PCT-US94-06264-3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             35 APLATELRCQCLQTLQGIHLKNIQSVKVKSPGPHCAQTEVIATLKNGQKACLNPASPMVK 94
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Patent No. 5844084
GENERAL INFORMATION:
APPLICANT: Guegler, Karl J.
APPLICANT: Hawkins, Phillip R.
APPLICANT: Wilde, Craig G.
APPLICANT: Wilde, Craig G.
APPLICANT: Sellhamer, Josef L.
TITLE OF INVENTION: INFLAMED ADENOID, ITS PRODUCTION AND USES
APPLICANT: Hawkins, Phillip R.
APPLICANT: Wilde, Craig G.
APPLICANT: Seilhamer, Jeffrey J.
TITLE OF INVENTION: A NOVEL CHEMOKINE EXPRESSED IN
TITLE OF INVENTION: INFLAMED ADENOID, ITS PRODUCTION AND USES
NUMBER OF SEQUENCES: 9
CORRESONDENCE ADDRESS:
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100.0%; Pred. No. 2.7e-40;
tive 0; Mismatches 0;
                                                                                                                                                                                        ADDRESSEE: Incyte Pharmaceuticals, Inc. STREET: 3174 Porter Drive CITY: Palo Alto.
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                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: FastSEQ Version 1.5
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/352,324A
FILLING DATE: 07-DEC-1994
CLASSIFICATION: 435
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STREET: 3174 Porter Drive CITY: Palo Alto .
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUBBER: 08/862,607
FILING DATE: 23-MAY-1997
APPLICATION NUBBER: 08/352,324
FILING DATE: 07-DEC-1994
ATTORREY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; Sequence 4, Application US/09203235; Patent No. 6071701
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ Version 1.5
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTAL.

ZIP: 94304
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
"TYPE: IBM Compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     33,954
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TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-855-0555
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                                                                 LENGTH: 107 amino acids
TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME: Luther, Barbara J
REGISTRATION NUMBER: 33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 107 amino acids TYPE: amino acid
                                                                                                                                                                                         Query Match
Best Local Similarity luv...
73; Conservative
                TELEX: N/A
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
512/474-7477
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        415-852-0195
                                                                                                                                   ; MOLECULE TYPE: peptide US-08-468-819-6
                                                                                                       single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                  95 KIIEKMLKNGKSN 107
                                                                                                                                                                                                                                                                                                                                                    61 KIIEKMLKNGKSN 73
                                                                                                                       linear
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                                                                                                                         TOPOLOGY:
TELEFAX:
TELEX: N
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Patent No. 5871723
GENERAL INFORMATION:
APPLICANT: Strieter, Robert M.
APPLICANT: Polverini, Peter J.
APPLICANT: Kunkel, Steven L.
TITLE OF INVENTION: CXC Chemokines as Regulators of TITLE OF SEQUENCES: 93
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDIUM TYPE: Floppy disk COMPUTER: IBM PC COMPATIble OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        100.0%; Score 379; DB 2; 100.0%; Pred. No. 2.7e-40;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ATTORNEY/AGENT INFORMATION:
NAME: Highlander, Steven L.
REGISTRATION NUMBER: 37,642
REFERENCE/DOCKET NUMBER: UMIC:003/HYL
                                                                                                                                                                                     NAME: Luther, Barbara J.
REGISTRATION NUMBER: 33,954
REFERENCE/DOCKET NUMBER: PF-0025 US
TELECOMUNICATION INFORMATION:
TELEPHONE: 415-855-0555
TELEFAX: 415-852-0195
INFORMATION FOR SEQ ID NO: 4:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US/08/468,819
FILING DATE: Concurrently herewith
CLASSIFICATION: 424
                                              CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/862,607
FILING DATE: 23-MAY-1997
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         E: Arnold, White & Durkee P.O. Box 4433
                                                                                                                  PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/352,324
FILING DATE: 07-DEC-1994
ATTORNEY/AGENT INFORMATION:
                   SYSTEM: DOS
FastSEQ Version 1.5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELECOMMUNICATION INFORMATION:
IBM Compatible
                                                                                                                                                                                                                                                                                                               SEQUENCE CHARACTERISTICS:
LENGTH: 107 amino acids
TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              512/418-3000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match 100.
Best Local Similarity 100.
Matches 73; Conservative
                                                                                                                                                                                                                                                                                                                                                                                               MOLECULE TYPE: peptide US-08-862-607-4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COUNTRY: US
ZIP: 77210
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   61 KIIEKMLKNGKSN 73
                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY: linear
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                                   SOFTWARE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 10
US-08-468-819-6
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Gaps
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APPLICANT: Guegler, Karl J.

APPLICANT: Guegler, Phillip R.

APPLICANT: Wilde, Craig G.

APPLICANT: Seilhamer, Jeffrey J.

APPLICANT: Seilhamer, Jeffrey J.

TITLE OF INVENTION: A NOVEL CHEMOKINE EXPRESSED IN

TITLE OF INVENTION: INFLAMED ADENOID, ITS PRODUCTION AND USES

NUMBER OF SEQUENCES: 9

CORRESPONDENCE ADDRESS:
Length 107;
                                            Indels
                                            ö
100.0%; Score 379; DB 2;
100.0%; Pred. No. 2.7e-40;
iive 0; Mismatches 0;
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1 APLATELRCQCLQTLQGIHLKNIQSVKVKSPGPHCAQTEVIATLKNGQKACLNPASPMVK 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 3. Application US/08330163

Patent No. 5656724

GENERAL INFORMATION:
APPLICANT: Daly, Thomas J.
TITLE OF INVENTION: Chemokine-Like Proteins and Methods of TITLE OF INVENTION: USE
NUMBER OF SEQUENCES: 46
CORRESPONDENCE ADDRESS:
ADDRESSE: 46
STREET: 225 Franklin Street
CITY: Boston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 3, Application US/08482111
Patent No. 5789539
GENERAL INFORMATION:
APPLICANT: Daly, Thomas J.
APPLICANT: LaRosa, Gregory J.
TITLE OF INVENTION: Chemokine-Like Proteins and Methods of TITLE OF INVENTION: Use
NUMBER OF SEQUENCES: 70
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 73;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30B
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/330,163
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 367; DB 1; I
Pred. No. 5.4e-39;
0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: Fasse, J. Peter
REGISTRATION NUMBER: 32,983
REFRENCE/DOCKET NUMBER: 00231
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 542-5070
INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ö
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Best Local Similarity 97.3%;
Matches 71; Conservative
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ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                             05-AUG-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                73 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY: U.S.A.
ZIP: 02110-2804
COMPUTER READABLE FORM:
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02110-2804
                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILING DATE: 05
CLASSIFICATION:
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US-08-330-163-3
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                                                                                                                   1 APLATELRCQCLQTLQGIHLKNIQSVKVKSPGPHCAQTEVIATLKNGQKACLNPASPMVK 60
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GENERAL INFORMATION:
APPLICANT: INCYTE PHARMACEUTICALS, INC.
TITLE OF INVENTION: ANOVEL CHEMOKINE EXPRESSED IN INFLAMED
TITLE OF INVENTION: ADENOID, ITS PRODUCTION AND USES
NUMBER OF SEQUENCES: 9
CORRESPONDENCE ADDRESS:
ADDRESSEE: INCYTE PHARMACEUTICALS, INC.
STREET: 3174 Porter Drive
CITY: Palo Alto
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 107;
                                       Length 107;
                                                                           Indels
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Pred. No. 2.7e-40;
                                     Query Match 100.0%; Score 379; DB 3; Best Local Similarity 100.0%; Pred. No. 2.7e-40; Matches 73; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        REGISTRATION NUMBER: 33954
REFERENCE/DOCKET NUMBER: PF-0025 PCT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WordPefect 6.1/MS-DOS 6.2
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: PCT/US95/16144
FILING DATE: 07-DEC-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/352,324
FILLING DATE: 07-DEC-1994
ATTORNEY/AGENT INFORMATION:
NAME: LUTHER, BARBARA J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             .;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-855-0555
TELEFAX: 415-852-0195
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   107 amino acids
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MOLECULE TYPE: protein
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 95 KIIEKMLKNGKSN 107
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                                                                                                                                                                                             61 KIIEKMLKNGKSN 73
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Best Local Similarity
Matches 73; Conserv
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COUNTRY:

PCT-US95-16144-4

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TOPOLOGY:

RESULT 13

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 APLATELRCQCLQTLQGIHLKNIQSVKVKSPGPHCAQTEVIATLKNGQKACLNPASPMVK 60
                                                     CURRENT APPLICATION DATA:

CURRENT APPLICATION DATA:

APPLICATION DATA:

FILING DATE:

CLASSIFICATION WUMBER:

NAME: Fasse, J. Peter

REGISTRATION WUMBER: 32,983

REFERENCE/DOCKET NUMBER: 32,983

RELEPHONE: (617) 542-8906

INFORMATION FOR SEQ ID NO: 3:

SEQUENCE CHARACTERISTICS:

LENGTH: 73 amino acids

STRUEL ANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 367; DB 1; L/
Pred. No. 5.4e-39;
0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 15
US-07-778-413E-18
Sequence 18, Application US/07778413E
Setent No. 5401651
GENERAL INFORMATION:
TITLE OF INVENTION: No. 5401651e1 Neutrophil
TITLE OF INVENTION: Activating Factors
NUMBER OF SEQUENCES: 22
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.5 in., DS, 2.0 MB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ATTORNEY AGENT INFORMATION:
NAME: COOK, ROBERT R.
REGISTRATION NUMBER: 31602
RECENENCE DOCKET NUMBER: A-204
TELECOMMUNICATION INFORMATION:
TELEPHONE: (805) 499-8011
INFORMATION FOR SEQ ID NO: 18:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MADLUM ITE.

COMPUTER: Apple Macintosh
OPERATING SYSTEM: Macintosh OS 7.0
SOFTWARE: Microsoft Word Version 5.1a
CURRENT APPLICATION DATA:
FILING DATE: 16-OCT-1991
CLASSIFICATION: 536
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADDRESSEE: Amgen Inc.
STREET: Amgen Center
STREET: 1840 Dehavilland Drive
CITY: Thousand Oaks
STATE: California
COUNTRY: USA
ZIP: 91320-1789
                 COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Ouery Match 96.8%;
Best Local Similarity 97.3%;
Matches 71; Conservative (
Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                    TOPOLOGY: linear
MOLECULE TYPE: peptide
US-08-482-111-3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             61 KIIEKMEKNGKSN 73
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GenCore version 4.5
Copyright (c) 1993 - 2000 Compugen Ltd.
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 protein search, using sw model OM protein

Run on:

May 4, 2001, 07:35:28; Search time 14.16 Seconds

(without alignments)
354.292 Million cell updates/sec

US-09-467-160-3 379 Title: Perfect score:

1 APLATELRCQCLQTLQGIHL......PASPMVKKIIEKMLKNGKSN Sequence:

Scoring table:

BLOSUM62 Gapop 10.0 , Gapext 0.5

198801 seqs, 68722935 residues rched: 198801 Total number of hits satisfying chosen parameters:

Minimum DB seq length: 0 Maximum DB seq length: 200000000

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

PIR_67:* Database :

pirl:* pir2:* pir3:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

	Description		macrophage inflamm	melanoma growth-st	GRO-gamma precurso	growth-regulated p	neutrophil chemo-a	gro-alpha precurso	macrophage inflamm	macrophage inflamm	macrophage inflamm	cytokine-induced n	gene KC protein -	alveolar macrophag	platelet basic pro	beta-thromboglobul	granulocyte chemot	neutrophil-activat	macrophage inflamm	transformation-ind	RSV-induced protei	granulocyte chemot	platelet factor 4	interferon gamma-i	platelet factor 4	cytokine - rabbit	interleukin-8 prec	C-X-C chemokine LI	platelet factor 4	platelet factor 4,	interleukin 8 - sh
		-																													
	QĮ.		JH0281	A28414	B38290	B28414	JN0572	A32954	JH0200	S21467	155614	S46198	164831	B44253	S42881	TGHU	B54188	A55010	151886	A26736	150417	A54188	S 69133	JN0470	A26774	S17507	A37034	A57325	PFHU4	A35766	S42496
	DB	:	7	~	7	~	7	~	~	7	7	7	~	~	7	٦	~	7	~	7	7	~	7	7	7	7	7	7	٦	7	7
	Length		107	107	107	101	96	96	100	100	100	100	53	117	119	128	75	114	53	103	103	75	90	125	105	36	66	132	101	126	101
dip	Query		100.0	87.3	85.8	•	74.9	73.4	68.8	65.7	65.7	61.5	55.9	55.4	55.4	55.1	54.9	52.6	47.0	46.6	46.6	•	45.6	•	43.9	43.8	43.4		٠	42.2	41.0
	Score		379	331	325	304	284	278	261	249	249	233	212	210	210	209	208	199.5	178	176.5	176.5	173	173	168	166.5	166	164.5	163	160	50	155.5
į	Result No.		1	2	е	7	S	9	7	80	σ	10	11	12	13	14	15	16	17	18	. 19	20	21	22	~	. 24	25	56	27	28	53

interleukin-8 - sh	cytokine boGRO - b	platelet factor 4	interleukin-8 - ra	alveolar macrophag	interleukin-8 prec	interleukin-8 - do	platelet factor 4	Neutrophil attract	interferon gamma-i	lipopolysaccharide	IP-10 precursor -	Mob-1 - rat	chemokine BRAK - h	pre-B-cell growth-	interleukin-8 homo	
146997	S50035	PFB04	146871	A44253	A53096	JN0841	PFHU4A	148148	TGHUGI	S33290	A45492	159277	JG0182	A53497	153416	
7	7	٦	7	7	7	7	7	7	٦	7	7	~	7	7	7	
101	33	88	101	103	103	95	104	101	86	40	86	96	66	88	8	
41.0	39.8	39.8	39.7	38.7	38.7	38.4	37.5	34.4	29.8	28.2	27.7	27.7	23.1	21.5	21.5	
155.5	151	151	150.5	146.5	146.5	145.5	142	130.5	113	107	105	105	87.5	81.5	81.5	
30	31	32	33	34	35	36	37	38	39	40	41	42	43	44	45	

ALIGNMENTS

	KESULT 1 JH0281 macrochade inflammatory protein 2 alpha prequireor - human
	N. Alternate names: gro-beta; growth required protein beta; melanoma growth-stimulato
	C; Species: Homo sapiens (man)
	C;Date: 30-Sep-1991 #sequence_revision 30-Sep-1991 #text_change 20-Aug-1999
	C; Accession: JH0281; A35931; A38290; A60407
	R; Tekamp-Olson, P.; Gallegos, C.; Bauer, D.; McClain, J.; Sherry, B.; Fabre, M.; van
	J. Exp. Med. 172, 911-919, 1990
	A; Title: Cloning and characterization of cDNAs for murine macrophage inflammatory pro
	A; Reference number: JH0200; MUID:90354792
	A; Accession: JH0281
	A; Molecule type: mRNA
	A; Residues: 1-107 <tek></tek>
	A; Cross-references: GB:X53799; NID:g34658; PIDN:CAA37808.1; PID:g34659
	R; Iida, N.; Grotendorst, G.R.
-	Mol. Cell. Biol. 10, 5596-5599, 1990

MACL Cell. Biol. 10, 5596-5599, 1990
MyTitle: Cloning and sequencing of a new gro transcript from activated human monocyte A; Title: Cloning and sequencing of a new gro transcript from activated human monocyte A; Reference number: A35931; MyID:90377259
A; Molecule type: mRNA
A; Molecule type: mRNA
A; Residues: 1-107 < LID>
A; Cross-references: GB:M57731; GB:M36964; NID:9183626; PIDN:AAA63182.1; PID:9183627
A; Cross-references: GB:M57731; GB:M36964; NID:9183626; PIDN:AAA63182.1; PID:9183627
B; Haskill, S.; Peace, A.; Morris, J.; Sporn, S.A.; Anisowicz, A.; Lee, S.W.; Smith, T.
Broc. Natl. Acad. Sci. U.S.A. 87, 7732-7736, 1990
A; Title: Identification of three related human GRO genes encoding cytokine functions. A; Reference number: A38290; MyID:91017578

A; Molecule type: mRNA A; Residues: 1-107 <HAS>

A.Cross-references: GB:M36820; NID:g183628; PIDN:AAA63183.1; PID:g183629
R.Sporn, S.A.; Eterman, D.F.; Johnson, C.E.; Morris, J.; Martin, G.; Ladner, M.; Hask J. Immunol. 144, 4434-4441, 1990
A.Title: Monocyte adherence results in selective induction of novel genes sharing hom A.Reference number: A60407; MUID:90257367
A.Accession: A60407

A; Status: not compared with conceptual translation

C; Superfamily: beta-thromboglobulin A; Molecule type: mRNA A; Residues: 56-107 <SPO>

C; Reywords: inflammation ... F;l-34/Domain: signal sequence #status predicted <SIG>F;l-34/Domain: signal sequence #status protein 2 alpha #status predicted <MAT>

Gaps ö Length 107; 0; Indels 100.0%; Score 379; DB 2; 100.0%; Pred. No. 1.3e-35; tive 0; Mismatches 0; Query Match 100.0 Best Local Similarity 100.0 Matches 73; Conservative

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us-09-467-160-3.rpr

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A; Molecule type: mRNA
A;Residues: 1-26,'G',29-107 <HAS>
A;Cross-references: GB:M3681; NID:g183632; PIDN:AAA63184.1; PID:g183633
A;Cross-references: GB:M3681; NID:g183632; PIDN:AAA63184.1; PID:g183633
B;Proost, P.; De Wolf-Peeters, C.; Conings, R.; Opdenakker, G.; Billiau, A.; Van Damm
J: Immunol. 150, 1000-1010, 1993.
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C;Species: 11.May-1991 #sequence_revision 27-Oct-1995 #text_change 20-Aug-1999
C;Accession: JH0282; B38290; C46519
R;Tekamp-Olson, P.; Gallegos, C.; Bauer, D.; McClain, J.; Sherry, B.; Fabre, M.; van J. Exp. Med. 172, 911-919, 1990
A;Title: Cloning and characterization of cDNAs for murine macrophage inflammatory pro A;Reference number: JH0200; MUID:90354792
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A;Accession: C46519
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R;Haskill, S.; Peace, A.; Morris, J.; Sporn, S.A.; Anisowicz, A.; Lee, S.W.; Smith, T Proc. Natl. Acad. Sci. U.S.A. 87, 7732-7736, 1990
A;Title: Identification of three related human GRO genes encoding cytokine functions. A;Reference number: A38290; MUID:91017578
A;Map position: 4q21-4q21
C;Superfamily: beta-thromboglobulin
F;l-34/Domain: signal sequence #status predicted <SIG>
F;35-107/Product: melanoma growth-stimulatory activity #status experimental <MAT>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  N; Alternate names: growth-regulated protein gamma; macrophage inflammatory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 APLATELRCQCLQTLQGIHLKNIQSVKVKSPGPHCAQTEVIATLKNGQKACLNPASPMVK 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 APLATELRCQCLQTLQGIHLKNIQSVKVKSPGPHCAQTEVIATLKNGQKACLNPASPMVK 60
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F;1-34/Domain: signal sequence #status predicted <SIG>
F;35-107/Product: GRO-gamma #status experimental <MAT>
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                                                                                                                                                                                                                                                                                                                                                  Score 331; DB 2;
Pred. No. 3.2e-30;
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Pred. No. 1.5e-29;
6; Mismatches 6
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A; Experimental source: MG-63 osteosarcoma cells
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87.7%;
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Matches 61; Conservative
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95 KIIEKMLNSDKSN 107
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KIIEKILNKGSTN 107
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A; Residues: 1-107 <TEK>
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A;Residues: 35-63, X',65 <SC2>
A;Experimental source: LPS-simulated monocytes
R;Experimental source: LPS-simulated monocytes
A;Experimental source: LPS-simulated monocytes
A;Reference number: A46519; MUID:93139489
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A; Residues: 35-42, 'X', 44, 'X', 46-48 <SCH>
A; Experimental source: dermal fibroblasts
R; Golds, E.E.; Mason, P.; Nyirkos, P.
Biochem. J. 259, 585-588, 1989
A; Fitle: Inflammatory cytokines induce synthesis and secretion of 9ro protein and a neut
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A;Reference number: A47626; MUID:90217938
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A;Reference number: S00983; MUID:88328991
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R;Schroeder, J.M.; Sticherling, M.; Henneicke, H.H.; Preissner, W.C.; Christophers, E.
J Immunol. 144, 223-2232, 1990
A;Title: IL-lalpha or tumor necrosis factor-alpha stimulate release of three NAP-1/IL-8-A;Reference number: A60401; MUID:90187866
A;Accession: B60401
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Diccule type: protein
A; Redidues: 35-41, X', 43-49, X', 51-52, XX', 55-57 <GOL>
R; Schroeder, J.M; Persoon, N.L.M.; Christophers, E.
J. Exp. Med. 171, 1091-1100, 1990
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A;Cross-references: GDB:120181; OMIM:155730
                                                                                                        95 KIIEKMLKNGKSN 107
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                                                    61 KIIEKMLKNGKSN
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A; Residues: 1-107 <BAK>
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A; Molecule type: mRNA
A; Residues: 1-107 <RIC>
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1 APLATELRCQCLQTLQGIHLKNIQSVKVKSPGPHCAQTEVIATLKNGQKACLNPASPMVK 60
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       A;Status: preliminary
A;Molecule type: protein
A;Residues: 25-57 <NA2>
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Best Local Similarity
Matches 50; Conserv
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A; Molecule type: mRNA
A; Residues: 1-96 <RYS>
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85 KIVQKMLK 92
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Nalternate names: CINC; cytokine-induced neutrophil chemoattractant; interleukin-8-like C; species: Rattus norvegitus (Morway rat) (C; becies: Rattus norvegitus (Morway rat) (C; Date: 30-Sep-1993 #sequence_revision 30-Sep-1993 #text_change 20-Jun-2000 C; Date: 30-Sep-1993 #sequence_revision 30-Sep-1993 #text_change 20-Jun-2000 C; Accession: JN0572; JQ1519; A34481; A48988; B48988; S51214 R; Konishi, K.; Takata, Y.; Yamamoto, M.; Yomogida, K.; Watanabe, K.; Tsurufuji, S.; Fuji Gene 126, 285-286, 1993 A; A; Athle: Structure of the gene encoding rat neutrophil chemo-attractant Gro. A; Reference number: JN0572; MUID:93246259
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A.Residues: 1-32,'S',34-96 <HUA>
A.Cross-references: GB.M86536
A.Experimental source: alveolar macrophage
A.Pxperimental source: alveolar macrophage
A.Note: the authors translated the codon AGT for residue 33 as Cys, AAC for residue 46 a
R.Watanabe, K.; Konishi, K.; Kujioka, M.; Kinoshita, S.; Nakagawa, H.
B. Biol. Chem. 264, 1955-19563, 1989
A.Fitle: The neutrophil chemoattractant produced by the rat kidney epithelioid cell line
A.Reference number: A34481; MUID:90062049
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A; Molecule type: protein
A; Molecule type: protein
A; Residues: 25-96 <WAT>
A; Residues: 25-96 <WAT>
A; Residues: 25-96 <WATA
B; Nakagawa, H.; Ikesu, A.; Hatakeyama, S.; Kato, H.; Gotoda, T.; Komorita, N.; Watanabe Biochem. Pharmacol. 45, 1425-1430, 1993
A; Title: Production of an interleukin-8-11ke chemokine by cytokine-stimulated rat NRK-49
                                                                                                                                                                              in transformed Chinese
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A. Residues: 1-96 <KON>
A. Residues: 1-96 <KON>
A. Cross references: DDBJ:D11445; NID:g391854; PIDN:BAA02009.1; PID:g220755
B. Huang, S.; Paulauskis, J.D.; Koblik, L.
Cochem. Biophys. Res. Commun. 184, 922-929, 1992
C. Title: Rat KC cDNA cloning and mRNA expression in lung macrophages and fibroblasts.
A. Reference number: J01519; MUID:92246987
A. A. Recession: J01519
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C.Species: Cricetulus griseus (Chinese hamster)
C.Date: 30-Jun_1989 #sequence_revision 30-Jun-1989 #text_change 20-Aug-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 APLATELRCQCLQTLQGIHLKNIQSVKVKSPGPHCAQTEVIATLKNGQKACLNPASPMVK 60
                                                                                                                                                                                                                                                                              A;Molecule type: mRNA
A;Residues: 1-101 <aN1>
A;Cross-references: GB:J03560; NID:g191088; PIDN:AAA36985_1; PID:g304509
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                                                                                                                                                                                                                                                                                                                                                                                                                  C;Superfamily: beta-thromboglobulin
F;1-23/Domain: signal sequence #status predicted <SIG>
F;24-101/Product: growth-regulated protein #status predicted <MAT>
                                                                                         R;Anisowicz, A.; Bardwell, L.; Sager, R.
Proc. Natl. Acad. Sci. U.S.A. 84, 7188-7192, 1987
A;Title: Constitutive overexpression of a growth-regulated gene.
A;Reference number: A94184; MUID:88041072
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 101;
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A;Residues: 25-57 <NAK>
A;Experimental source: kidney, NRK-49F fibroblasts
A;Note: sequence extracted from NCBI backbone (NCBIP:129132)
A;Accession: B48988
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match 80.2%; Score 304; DB 2; L
Best Local Similarity 77.1%; Pred. No. 3.3e-27;
Matches 54; Conservative 10; Mismatches 6;
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A; Accession: A48988
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                                                                   C; Accession: B28414
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A.Experimental source: kidney, NRK-49F fibroblasts
A.Note: sequence extracted from NCBI backbone (NCBIP:129131)
R.Hanzawa, H.; Haruyama, H.; Watanabe, K.; Tsurufuji, S.
REBS Lett. 354, 207-212, 1994
A.FIILE: The three dimensional structure of rat cytokine CINC/Gro in solution by homo A.Reference number: 551214; MUID:95046335
A.Contents: annotation; conformation by (1)H-NMR, residues 25-96
A.Accession: S51214
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C; Species: Mus musculus (house mouse)
C; Date: 20-Oct-1989 #sequence_revision 20-Oct-1989 #text_change 20-Aug-1999
C; Accession: A32954; JH0081
R; Oquendo, P.; Alberta, J.; Wen, D.; Graycar, J.L.; Derynck, R.; Stiles, C.D.
A; Title: The platelet-derived growth factor-inducible KC gene encodes a secretory pro A; Reference number: A32954; MUID:89139485
A; Residues: 1-96 <OQU>
                                                                                                                                                                                                                                                                                              A;Molecule type: protein
A;Residues: 25-96 <HAN>
C;Comment: This protein has chemotactic activity for neutrophils and has melanoma gro
C;Genetics:
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R:Ryseck, R.P.: MacDonald-Bravo, H.; Mattel, M.G.; Bravo, R.
Exp. Cell Res. 180, 266-275, 1989
A;Title: Cloning and sequence of a secretory protein induced by growth factors in mou A;Reference number: JH0081; MUID:89078502
                                                                                                                                                                                                                                                                                                                                                                                                                                       A.Gene: gro; KC
A.Introns: 24/1; 65/2; 92/2
C.Superfamily: beta-thromboglobulin
C.Keywords: cytokine; disulfide bond
F.1-24/Domain: signal sequence #status predicted <SIG>
F.25-96/Product: neutrophil chemo-attractant Gro protein #status experimental <CYT>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        25 APVANELRCQCLQTVAGIHFKNIQSLKVMPPGPHCTQTEVIATLKNGREACLDPEAPMVQ 84
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73.5%; Pred. No. 2.6e-24;
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75.0%; Pred. No. 5.4e-25;
Live 10; Mismatches 7;
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C;Reywords: extracellular protein
F:1-24/Domain: signal sequence #status predicted </F;25-96/Product: gro-alpha #status predicted <MAT>
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Matches 51; Conservative
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61 85

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C;Species: Rattus norvegicus (Norway rat)
C;Species: S46198; C48988
R;Nakagawa, H.; Komorita, N.; Shibata, F.; Ikesue, A.; Konishi, K.; Fujioka, M.; Kato Biochem. J. 301, 545-550, 1994
A;Title: Identification of cytokine-induced neutrophil chemoattractants (CINC), rat Gences and characterization.
A;Reference number: S46198; MUID:94318061
A;Status: preliminary
A;Status: preliminary
A;Status: preliminary
A;Status: Residues: 1-100 < NARX
A;Residues: 1-
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C;Date: 26-Jul-1996 #sequence_revision 26-Jul-1996 #text_change 20-Aug-1999
C;Date: 26-Jul-1996 #sequence_revision 26-Jul-1996 #text_change 20-Aug-1999
C;Accession: I55614
R;Feng, L.; Xia, Y.; Yoshimura, T.; Wilson, C.B.
A;Title: Modulation of neutrophil influx in glomerulonephritis in the rat with anti-m A;Reference number: I55614; WUID:95189993
                                                                                                                                                                                                                                                                                                 30 VASELRCQCLTTLPRVDFKNIQSLTVTPPGPHCAQTEVIATLKDGHEVCLNPEAPLVQRI 89
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
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C;Superfamily: beta-thromboglobulin
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                                                                                 ; Score 249; DB 2; Length 10; Pred. No. 4.8e-21; 14; Mismatches 13; Indels
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A:Note: sequence extracted from NCBI backbone (NCBIP:129130)
C;Superfamily: beta-thromboglobulin
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                                                                                      65.7%;
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90 VQKILNKGKAN 100
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A; Status: preliminary
A; Molecule type: protein
A; Residues: 33-56 <NA2>
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                                                                                      Query Match
Best Local Similarity
Matches 44; Conserv
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A; Residues: 1-100 <RES>
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A.Status: preliminary
A.Molecule type: mRNA
A.Fosidues: 1-100 CADRIA
A.Fosidues: 1-100 CADRIA
A.Cross references: EMBL.X65647; NID:956665; PIDN:CAA46599.1; PID:956666
R.Nakagawa, H.; Ikesue, A.; Hatakeyama, S.; Kato, H.; Gotoda, T.; Komorita, N.; Watanabe
Blochem. Pharmacol. 45, 1425-1430, 1993
A.Patile: Production of an interleukin-8-like chemokine by cytokine-stimulated rat NRK-49
A.Facession: D48988
A.Facession: D48988
A.Status: proliminary
A.Molecule type: protein
A.Fasidues: 32-45 cNAK>
A.Fasidues: 32-45 cNAK>
A.Fasidues: 32-45 cNAK>
A.Fasidues: 32-45 cNAK>
A.Fasidues: Sequence extracted from NCBI backbone (NCBIP:129129)
                                                                                                                                                                                                                                                                                                                          JH0200

macrophage inflammatory protein 2 precursor - mouse
C;Species 10.
C;Species 10.
C;Date: 30-Sep-1991 #sequence_revision 30-Sep-1991 #text_change 20-Aug-1999
C;Accession: JH0200; A32190
C;Accession: JH0200; A32190
C;Accession: JH0200; A32190
C;Accession: JH0200; A3190
C;Batlegos, C.; Bauer, D.; McClain, J.; Sherry, B.; Fabre, M.; van Dev A:Title: Choing and characterization of cDNAs for murine macrophage inflammatory protein A:Reference number: JH0200; MUID:90354792
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A; Cross-references: GB: X53798; NID:g53128; PIDN:CAA37807.1; PID:g53129
Wolpe, S.D.; Sherry, B.; Juers, D.; Davatelis, G.; Yurt, R.W.; Cerami, A.
Oc. Natl. Acad. Sci. U.S.A. 86, 612-616, 1989
W; Title: Identification and characterization of macrophage inflammatory protein 2.
A; Reference number: A32190; MUID:89098980
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A.Resiques: 28-59 <WOL>
C.Superfamily: beta-thromboglobulin
C.Superfamily: beta-thromboglobulin
C.Keywords: heparin binding
F:1-27/Domain: signal sequence #status predicted <SIG>
F:28-100/Product: macrophage inflammatory protein 2 #status experimental <MAT>
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C;Date: 20-Feb-1995 #sequence_revision 20-Feb-1995 #text_change 20-Aug-1999
C;Accession: S21467; D48988
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               28 AVVASELRCQCLKTLPRVDFKNIQSLSVTPPGPHCAQTEVIATLKGGQKVCLDPEAPLVQ 87
25 APIANELRCQCLQTMAGIHLKNIQSLKVLPSGPHCTQTEVIATLKNGREACLDPEAPLVQ 84
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           68.9%; Score 261; DB 2;
65.8%; Pred. No. 2.2e-22;
Live 12; Mismatches 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Ribriscoll, K. submitted to the EMBL Data Library, April 1992 M. Reference number: $21467 A; Accession: $21467
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            crophage inflammatory protein 2 - rat
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Matches 48; Conservative
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88 KIIQKILNKGKAN 100
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A; Molecule type: mRNA
A; Residues: 1-100 <TEK>
                                                                                 KIIEKMIK 68
                                                                                                                             ||::||||
KIVQKMLK 92
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Query Match

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A.Title: Characterization of human platelet basic protein, a precursor form of low-af A.Reference number: A2448; MUID:86216117
A.Accession: A24448
A.Accession: D.A.Accession: D.A.Accession: D.A.Accession: D.A.Accession: D.A.Accession: D.A.Accession: PLO222
A.Accession: D.A.Accession: D.A.Ac
                                                                                                                                                                                                          platelet basic protein - pig
C;Species: Sus scrofa domestica (domestic pig)
C;Date: 06-Jan-1995 #sequence_revision 06-Jan-1995 #text_change 20-Aug-1999
C;Accession: S43460; S42881
R;Power, C.A.; Proudfoot, A.E.I.; Magnenat, E.; Bacon, K.B.; Wells, T.N.C.
Eur. J. Biochem. 221, 713-719, 1994
A;Title: Molecular cloning and characterisation of a neutrophil chemotactic protein f
A;Reference number: S43460; MUID:94229068
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               N.Contains: connective-tissue activating peptide III; CTAP-III; histamine-releasing f C; Species: Homo sapiens (man)
C; Species: Homo sapiens (man)
C; Date: 30-Apr-1979 #sequence_revision 31-Dec-1992 #text_change 08-Dec-2000
C; Accession: A39546; A37382; A24448; PL0222; A3982; A90411; A60709; A61240; B61240; R.Majumdar, S.; Gonder, D.; Kouteis, B.; Poncz, M.
J. Biol. Chem. 266, 5785-5789, 1991
A; Title: Characterization of the human beta-thromboglobulin gene. Comparison with the A; Reference number: A39546; MUID:91170256
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A;Residues: 1-119 <POW>
A;Cross-references: EMBL:X77935; NID:g457753; PIDN:CAA54907.1; PID:g457754
C;Superfamily: beta-thromboglobulin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             51 ELRCLCLNTVSGIHPSNIQSLEVIRAGAHCAKVEVIATLKNDKKICLDPEAPRIKKIVQK 110
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A; Residues: 1-128 < MAJ>
A; Residues: 1-128 < MAJ>
A; Cross-references: GB:M54995; NID:g181175; PIDN:AAA62836.1; PID:g181176
A; Note: the authors translated the codon GAT for residue 109 as Pro
R; Wenger, R.H.; Wicki, A.N.; Walz, A.; Kieffer, N.; Clemetson, K.J.
Blood 73, 1498-1503, 1988-1503 of connective tissue activating peptide
A; Fitle: Cloning of cDNA coding for connective tissue activating peptide
A; Reference number: A37382; MUID:89229374
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                beta-thromboglobulin precursor [validated] - human
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Matches 38; Conservative
104 KIVQKMLDSGK 114
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A; Molecule type: mRNA
A; Residues: 1-128 < WEN>
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R; Huang, S.; Paulauskis, J.D.; Godleski, J.J.; Kobzik, L.
A; Huang, S.; Paulauskis, J.D.; Godleski, J.J.; Kobzik, L.
A; Title: Expression of macrophage inflammatory protein-2 and KC mRNA in pulmonary inflam A; Reference number: 151886; MUID:93035653
A; Accession: 164831
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R; Goodman, R.B.; Foster, D.C.; Mathewes, S.L.; Osborn, S.G.; Kuijper, J.L.; Forstrom, J. Biochemistry 31, 10483-10490, 1992
A; Title: Wolecular cloning of porcine alveolar macrophage-derived neutrophil chemotactic A; Reference number: A44253; MUID: 93041741
A; Accession: B44253
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c;Species: Sus scrofa domestica (domestic pig)
C;Date: 30-Apr-1993 #sequence_revision 18-Nov-1994 #text_change 20-Aug-1999
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Species: Rattus sp. (rat)
Date: 02-Aug-1996 #sequence_revision 02-Aug-1996 #text_change 23-Feb-1997
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A)Experimental source: alveolar macrophage
A;Note: sequence extracted from NCBI backbone (NCBIN:117417, NCBIP:117418)
C;Superfamily: beta-thromboglobulin
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Pred. No. 3.5e-17;
9; Mismatches 6; Indels
                                                                                                                                         12; Indels
                                                             Score 233; DB 2;
Pred. No. 3e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A;Status: preliminary; translated from GB/EMBL/DDBJ
A;Molecule type: mRNA
A;Residues: 1-53 <RES>
                                                                                                                                         13; Mismatches
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C;Superfamily: beta-thromboglobulin
                                                             61.5%;
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71.7%;
                                                                                                                                         Conservative
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Best Local Similarity 71.7'
Matches 38; Conservative
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A;Residues: 1-117 <GOO>
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                                                                                                       Similarity
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Matches 38; Conserv
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91 IQKLLKS 97
                                                                                                                                         42;
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Best Local S
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Job time: 56 sec
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R;Castor, C.W.; Miller, J.W.; Walz, D.A.
Proc. Natl. Acad. Sci. U.S.A. 80, 765-769, 1983
A;Title: Structural and biological characteristics of connective tissue activating pepti A;Reference number: A93982; MUID:83144010
A;Reference number: A93982; MUID:83144010
A;Residues: 44-66:125-128 CGAS>
R;Begg, G.S.; Pepper, D.S.; Chesterman, C.N.; Morgan, F.J.
Bjochemistry 17, 173-174-174, 1978
B;Residues: 44-66:125-128 CGAS>
R;Reference number: A90411; MUID:78187279
A;Reference number: A90411; MUID:78187279
A;Rolecule type: protein
A;Residues: 48-128 CBEC>
R;Beaza, M.L.; Reddigari, S.R.; Kornfeld, D.; Ramani, N.; Smith, E.M.; Hossler, P.A.; Fi
J; Clin. Invest. 85, 1516-1521, 1990
A;Reference number: A60709; MUID:90237229
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A; Molecule type: protein
A; Residues: 59-62, 'X', 64-79 < KA2>
B; Residues: 59-62, 'X', 64-79 < KA2>
B; Rungl, A.J.; Machius, M.; Huber, R.; Schwer, C.; Lam, C.; Aschauer, H.; Ehn, G.; Lindl
FEBS. Lett. 347, 300-303, 1994
A; Title: Purification, crystallization and preliminary X-ray diffraction analysis of red
A; Reference number: $46247; MUID: 94307404
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A; Contents: annotation c; Comment: There appears to be a second beta-thromboglobulin-like human gene. C; Comment: There appears to be a second beta-thromboglobulin-like human gene. C; Comment: Connective tissue activating peptides (CTAP) are a group of proteins capable latelet, and polymorphonuclear leukocyte origin, respectively. C; Comment: CTAP-III, a monomer isolated from platelets, stimulates DNA synthesis, mitosi & sulfated glycosaminoglycan. It also stimulates the formation and secretion of plasming C; Comment: Proteolytic removal of the first four residues of CTAP-III produces the active wed growth factor.
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C.Superfamily: beta-thromboglobulin
C.Superfamily: beta-thromboglobulin
C.Superfamily: beta-thromboglobulin
C.Superfamily: beta-thromboglobulin
T.34 Domain: signal sequence #status predicted <SIG>
T.54 Domain: propeptide #status predicted <PRO>
F.44 128 Product: connective-lissue activating peptide III #status experimental <CTAP>
F.59-128/Product: beta-thromboglobulin #status experimental <PRG>
F.59-128/Product: neutrophil-activating peptide 2 #status experimental <NAP2>
F:63-89,65-105/Disulfide bonds: #status experimental
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A.Accession: A60709
A.Molecule type: protein
A.Residues: 44-62, X', 64-79 < BAE>
Kaplan, A.P.; Baeza, M.; Reddigari, S.; Kuna, P.
L. Arch. Allergy Appl. Immunol. 94, 148-153, 1991
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A;Cross-references: GDB:127391; OMIM:121010
A;Map position: 4pl2-4q13
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A, Accession: A61240
A, Molecule type: protein
A, Residues: 44-61, /XX', 64 <KAP>
A, Accession: B61240
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Best Local Similarity 55.7
Matches 39; Conservative
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63 IEKMLKNGKS 72 ::| | :| 117 VQKKLAGDES 126

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RESULT 15 B54188

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granulocyte chemotactic protein, GCP-2 - bovine
C;Species: Bos primigenius taurus (cattle)
C;Species: Bos primigenius taurus (cattle)
C;Accession: B54188
R;Proost, P.; Wuyts, A.; Conings, R.; Lenaerts, J.P.; Billiau, A.; Opdenakker, G.; Va Biochmaistry 32, 10170-10177, 1993
A;Fitle: Human and bovine granulocyte chemotactic protein-2: complete amino acid sequence number: A54188; MUID: 94001982
A;Fitle: Human and bovine granulocyte chemotactic protein-2: complete amino acid sequence number: A54188; MUID: 94001982
A;Fitle: Human and bovine granulocyte chemotactic protein-2: complete amino acid sequence humber: A54188; MUID: 94001982
A;Fatus: preliminary
A;Molecule type: protein
A;Scatus: preliminary
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Fri May

GenCore version 4.5 Copyright (c) 1993 - 2000 Compugen Ltd.

OM protein - protein search, using sw model

May 4, 2001, 07:35:28 ; Search time 9.61 Seconds (without alignments) 260.213 Million cell updates/sec Run on:

US-09-467-160-3
379
1 APLATELRCQCLQTLQGIHL.....PASPMVKKIIEKMLKNGKSN 73 Title: Perfect score: Sequence:

BLOSUM62 Gapop 10.0 , Gapext 0.5 Scoring table:

93435 seqs, 34255486 residues arched:

93435 Total number of hits satisfying chosen parameters:

Minimum DB seq length: 0 Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

SwissProt_39:* Database : Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

	P19875 homo sapien P09341 homo sapien P09341 homo sapien P09341 cricetulus O46678 ovis arress O46675 bos taurus O46675 bos taurus O46675 bos taurus O46676 bos taurus O46677 bos taurus P130782 oryctolagus P14095 rattus norv P12850 mus musculu O55235 cavia porce P3034 rattus norv Q1074 rattus norv Q1074 rattus norv P12952 sus scrofa P02775 homo sapien P02775 homo sapien P08317 gallus gall P08317 gallus gall P08317 gallus gall P08317 pailus gallus gall P08317 homo sapien P08317 pailus gall P08315 homo sapien P04328 homo sapien P04328 mus morv P10145 hinterleuk P46653 cercocebus P50228 mus musur porve
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Length DB	1001 1007 1007 1008 1008 1008 1008 1008
% Query Match Le	1000 8877.0 8877.0 8877.0 8877.0 8877.0 887.0 87.0
Score	2000 2000 2000 2000 2000 2000 2000 200
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P02776 homo sapien P18340 mus musculu	P79255 bos taurus P36925 ovis aries P02777 bos taurus	P19874 oryctolagus P41324 canis famil	P26894 sus scrofa P10720 homo sapien	062812 equus cabal P49113 cavia porce	P02778 homo sapien
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MEDLINE-88328991; PubMed=2970963;
Richmond A., Balentien E., Thomas H.G., Flaggs G., Barton D.E.,
Spiess J., Bordoni R., Francke U., Derynck R.;
"Molecular characterization and chromosomal mapping of melanoma
growth stimulatory activity, a growth factor structurally related to
beta-thromboglobulin.",
EMBO J. 7:2025-2033(1988).
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PRINTS; PR00437; SMALLCYTKCXC.
PROSITE; PS00471; SMALL_CYTOKINES_CXC; 1.
Cytokine; Chemotaxis; Inflammatory response; Signal; 3D-structure.
SIGNAL 35 107 MACROPHAGE INFLAMMATORY PROTEIN-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     01-MAR-1989 (Rel. 10, Created)
01-MAR-1989 (Rel. 10, Last sequence update)
01-OCT-2000 (Rel. 40, Last annotation update)
GROWTH REGULATED PROTEIN PRECURSOR (MELANOWA GROWTH STIMULATORY
ACTIVITY) (MGSA) (NEDTROPHIL-ACTIVATING PROTEIN 3) (NAP-3).
SCYBI OR GROI OR GROA OR GRO OR MGSA.
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                                                                                                                                                                                                                                                                                                                                   Length 107;
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Anisowicz A., Bardwell L., Sager R.;
"Constitutive overexpression of a growth-regulated gene in
transformed Chinese hamster and human cells.";
Proc. Natl. Acad. Sci. U.S.A. 84:7188-7192(1987).
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                                                                                                                                                                                                                                                                                                 11389 MW; 740F277E928571BA CRC64;
                                                                                                                                                                                                                                                                                                                                 Score 379; DB 1;
Pred. No. 2.4e-36;
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                                                                          EMBL, X53799; CAA37808.1; -.
EMBL, M36820; AAA63183.1; -.
EMBL, M57731; AAA63182.1; -.
EMBL, AF043340; AAC03540.1; -.
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1 Similarity 100.0%;
73; Conservative 0
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                                                                                                                                                                                     PF00048; IL8; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                   61 KIIEKMLKNGKSN 73
                                                                                                                      PIR; JH0281; JH0281.
PDB; IQNK; 04-FEB-00.
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Best Local Similarity
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P09341;
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GRO_HUMAN
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SIMILARITY: BELONGS TO THE INTERCRINE ALPHA FAMILY (SMALL CYTOKINE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MEDLINE=89246368; PubMed=2655583; Golds E.E., Mason P., Nyirkos P.; "Inflammatory cytokines induce synthesis and secretion of gro protein and a neutrophil chemotactic factor but not beta 2-microglobulin in human synovial cells and fibroblasts."; Biochem. J. 259:585-588(1989).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Wen D., Rowland A., Derynck R.; ^{\rm R} ^{\rm R} ^{\rm R}
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J. Biol. Chem. 269:32809-32915(1994).
-!- FUNCTION: HAS CHEMOTACTIC ACTIVE FOR NEUTROPHILS. MAY PLAY A
ROLE IN INFLAMMATION AND EXERTS ITS EFFECTS ON ENDOTHELIAL CELLS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Schroeder J.-M., Persoon N.L.M., Christophers E.;
"Lipopolysaccharide-stimulated human monocytes secrete, apart from neutrophil-activating peptide l/interleukin 8, a second neutrophilactivating protein. NH2-terminal amino acid sequence identity with melanoma growth stimulatory activity.";
J. Exp. Med. 171:1091-1100(1990).
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Hl assignment and secondary structure determination of human
melanoma growth stimulating activity (MGSA) by NMR spectroscopy.";
FEBS Lett. 330:302-306(1993).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MEDLINE-94376296; PubMed-8089846; Fairbrother W.J., Reilly D., Colby T., Hesselgesser J., Horuk R.; "The solution structure of melanoma growth stimulating activity."; J. Mol. Biol. 242:255-270(1994).
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                                                                                                                                   Baker N.E., Kucera G., Richmond A.; "Nucleotide sequence of the human melanoma growth stimulatory
                                                                                                                                                                                                                                                               activity (MGSA) gene.";
Nucleic Acids Res. 18:6453-6453(1990).
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MEDLINE-89356650; PubMed-2670560;
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MEDLINE=93387459; PubMed=8397104;
                                                       MEDLINE=91057157; PubMed=2129556;
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PIR; S00983; S00983.
PIR; S03976; S03976.
PIR; S13669; S13669.
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97A69946B7F1F070 CRC64;
                                                                                                                                                                                                     Pfam; PP00048; IL8; 1.
PRINTS; PR00436; INTERLEUKIN8.
PRINTS; PR00437; SMALLCTYCCXC.
PROSITE; PS00471; SMALL_CYTOKINES_CXC; 1.
CYTOK ne; Chemotaxis; Inflammatory response; Signal.
SIGNAL
                                                                                                                                                                                                                                                                                                                                                                                                                                   85.8%; Score 325; DB 1; L6
83.6%; Pred. No. 3.2e-30;
"Mismatches 6;
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01-MAR-1989 (Rel. 10, Last sequence update)
01-OCT-2000 (Rel. 40, Last annotation update)
GROWTH REGULATED PROTEIN PRECURSOR.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               101 AA
                                                                                                                                                                                                                                                                                                                                             BY SIMILARITY.
BY SIMILARITY.
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                                                 EMBL, X53800; CAA37809.1; -.
EMBL, M36821; AAA63184.1; -.
PIR, B38290; B38290.
PIR, JH0282.
HSSP; P09341; IMGS.
MIM; 139111; -.
InterPro; IPR001891; -.
InterPro; IPR001811; -.
InterPro; IPR001811; -.
InterPro; IPR001811; -.
                                                                                                                                                                                                                                                                                                                                                                                                  11342 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  EMBL; J03560; AAA36985.1; -.
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85
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95 KIIEKILNKGSTN 107
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                                                                                                                                                                                                                                                                                                                                                                                                107 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE FROM N.A.
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NCBI_TaxID=10029;
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45
27
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P09340;
                                                                                                                                                                                                                                                                                                                                                             DISULFID
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                                                                                                                                                                                                                                                                                                                 CHAIN
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-!- FUNCTION: MAY PLAY A ROLE IN INFLAMMATION AND EXERT ITS EFFECTS
ON ENDOTHELIAL CELLS IN AN AUTOCRINE FASHION.
-!- SIMILARITY: BELONGS TO THE INTERCRINE ALPHA FAMILY (SMALL CYTOKINE
C-X-C) (CHEMOKINE CXC).
                                                                                                                                    PRINTS; PR00436; INTERLEUKINB.
PRINTS; PR00437; SMALLCYTKCXC.
PROSITE; PS00471; SMALL_CYTOKINES_CXC; 1.
Cytokine; Growth factor; Inflammatory response; Signal; 3D-structure.
                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       01-FEB-1991 (Rel. 17, Created)
01-FEB-1991 (Rel. 17, Last sequence update)
01-CT-2000 (Rel. 40, Last annotation update)
MACROPHAGE INFLAMMATORY PROTEIN -2-BETA PRECURSOR (MIP2-BETA) (GROWTH
REGULATED PROTEIN GAMMA) (GRO-GAMMA).
                                                                                                                                                                                                                                                                                                                                                                                                  1 APLATELRCQCLQTLQGIHLKNIQSVKVKSPGPHCAQTEVIATLKNGQKACLNPASPMVK 60
                                                                                                                                                                                                                                                                                                                                                                                                                  MEDLINE-91017578; PubMed-2217207;
Haskill S., Peace A., Morris J., Sporn S.A., Anisowicz A., Lee S.W.,
Smith T., Martin G., Ralph P., Sager R.;
"Identification of three related human GRO genes encoding cytokine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                                                                                ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TISSUE-Histiocytic lymphoma;
MEDLINE-90354792; PubMed=2201751;
Tekamp-Olson P., Gallegos C., Bauer D., McClain J., Sherry B.,
Fabre M., van Deventer S., Cerami A.;
"Cloning and characterization of cDNAs for murine macrophage inflammatory protein 2 and its human homologues.";
[2] Exp. Med. 172:911-919(1990).
                                                                                                                                                                                                                                                                                                                           Score 331; DB 1; Length 107;
Pred. No. 6.8e-31;
4; Mismatches 5; Indels
                                                                                                                                                                                                                                                                            11301 MW; 17048A6B4D765CA2 CRC64;
                                                                                                                                                                                                                          GROWTH REGULATED PROTEIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           107 AA.
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                                                                                                                                                                                                                                                                                                                             87.3%;
87.7%;
                                                                                                                                                                                                                                                                                                                                                              64; Conservative
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                                                               InterPro; IPR001089; -. InterPro; IPR001811; -. InterPro; IPR002473; -. Pfam; PF00048; IL8; 1.
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KIIEKMLNSDKSN 107
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     61 KIIEKMLKNGKSN 73
1MGS; 30-SEP-94.
1MSG; 31-MAR-95.
1MSH; 31-MAR-95.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens (Human).
                                                                                                                                                                                                                                                                          107 AA;
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                                                   MIM; 155730;
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Proc. Natl. /
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                                                                                                                                                                                                                                                          DISULFID
SEQUENCE
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Best Local
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Anisowicz A., Bardwell L., Sager R.;
Constitutive overexpression of a growth-regulated gene in transformed Chinese hamster and human cells.";
Proc. Natl. Acad. Sci. U.S.A. 84:7188-7192(1987).
-!- FUNCTION: HAS CHEMOTACITC ACTIVITY FOR NEUTROPHILS.
-!- SIMILARITY: BELONGS TO THE INTERCRINE ALPHA FAMILY (SMALL CYTOKINE C-X-C) (CHEMOKINE CXC).
                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 APLATELRCQCLQTLQGIHLKNIQSVKVKSPGPHCAQTEVIATLKNGQKACLNPASPMVK 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Cricetulus griseus (Chinese hamster).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Cricetinae;
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o
MACROPHAGE INFLAMMATORY PROTEIN-
                                                                                                                                                                                                                                                                                                                                     Length 107;
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CHAIN
DISULFID
DISULFID
SEQUENCE
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GROG_BOVIN
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                                                                                                                                                                                                                   ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE FROM N.A. Yoshimura T., Modi W.S.;
"Isolation of novel GRO genes, and a phylogenetic analysis of the CXC
                                                                                                                                                                                                                   0; Gaps
                                                                                                                                                                                                                                      1 APLATELRCQCLQTLQGIHLKNIQSVKVKSPGPHCAQTEVIATLKNGQKACLNPASPMVK 60
                                                                                                                                                                                                                                                    29 APVANELRCQCLQTMTGVHLKNIQSLKVTPPGPHCTQTEVIATIKNGQEACLNPEAPMYQ 88
                                                                                                                                                                                                                                                                                                                                                                                                                                           Ovis aries (Sheep).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora; Bovoidea;
                                                                                                                                                                                           80.2%; Score 304; DB 1; Length 101; 77.1%; Pred. No. 7.4e-28; ive 10; Mismatches 6; Indels
                                                                                                                                                                                                                 Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         InterPro: IPR001089; -
InterPro: IPR001081; -
InterPro: IPR001811; -
InterPro: IPR0043; -
Pfam; PF00048; ILB; 1.
PRINTS; PR00435; INTERLEUKINB.
PRINTS; PR00437; SMALLCYTKCXC.
PROSTTE; PS00471; SMALL_CYTOKINES_CXC; 1.
Cytokine; Growth factor; Inflammatory response; Signal.
SIGNAL
                 InterPro; IPR001089; -
InterPro; IPR001081; -
InterPro; IPR001811; -
InterPro; IPR00473; -
Pfan; PF00048; IL8; I.
PRINTS; PR00436; INTERLEUKIN8.
PRINTS; PR00437; SMALLCYTCXCC.

CYLOKINE; PG00471; SMALLCYTOKINES.CXC; 1.

CYLOKINE; Growth factor; Inflammatory response; Signal.
                                                                                                                       GROWTH REGULATED PROTEIN.
BY SIMILARITY.
BY SIMILARITY.
666FB7E9CC512019 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                  30-MAY-2000 (Rel. 39, Created)
30-MAY-2000 (Rel. 39, Last sequence update)
01-CCT-2000 (Rel. 40, Last annotation update)
GCOWTH REGULATED PROTEIN PRECURSOR.
SCYB1 OR GRO.
                                                                                                                                                                                                                                                                                                                                                                103 AA.
                                                                                                                POTENTIAL.
                                                                                                                                              39 79 E
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                                                                                                                                                                                        Query Match 80.2%;
Best Local Similarity 77.1%;
Matches 54; Conservative
                                                                                                                                                                                                                                                                                                                                                                STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Bovidae; Caprinae; Ovis.
                                                                                                            28
101
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                                                                                                                                                                                                                                                                                  61 KIIEKMLKNG 70
PIR; B28414; B2841
HSSP; P10889; 1MI2
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89 KIVQKMLKSG 98
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O46678;
                                                                                                                                  DISULFID
DISULFID
SEQUENCE
                                                                                                                SIGNAL
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YOShimura T., Modi W.S.;
Submitted (WAR-1997) to the EMBL/GenBank/DDBJ databases.
-!- SIMILARITY: BELONGS TO THE INTERCRINE ALPHA FAMILY (SMALL CYTOKINE C-x-C) (CHEMOKINE CXC).
                                                                                                                                                                                        Gaps
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                                                                                                                                                                                                                                                           1 APLATELRCQCLQTLQG1HLKNIQSVKVKSPGPHCAQTEVIATLKNGQKACLNPASPMVK 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GROWTH REGULATED PROTEIN HOMOLOG GAMMA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Bos taurus (Bovine).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora; Bovoidea;
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15-DEC-1998 (Rel. 37, Last sequence update)
15-DEC-1998 (Rel. 37, Last annotation update)
GROWTH REGULATED PROTEIN HOMOLOG GAMMA PRECURSOR (GRO-GAMMA).
                                                                                                                                           Length 103;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 98;
                                                                                                                                        80.2%; Score 304; DB 1; Length 10
76.7%; Pred. No. 7.5e-28;
tive 6; Mismatches 11; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 6; Indels
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InterPro; IPR001089; -.
InterPro; IPR001081; -.
Pfam; PF00048; ILS; 1.
PRINTS; PR00437; SMALLCYTKCXC.
PROSITE; PS00471; SMALL_CYTOKINES_CXC; 1.
CYtokine; Growth factor; Inflammatory response; Signal.
GROWTH REGULATED PROTEIN.
BY SIMILARITY.
BY SIMILARITY.
C59857F346716695 CRC64;
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83.6%; Pred. No. 2e-27;
live 5; Mismatches 6
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BY SIMILARITY
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103
65
81
10820 )
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91 KIIDKMLNQASSN 103
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                                                                                                                                                                                                                                                                                                                            61 KIIEKMLKNGKSN 73
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Best Local Similarity
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KIIDKML 97
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O46675;
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DISULFID
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GRO1_RABIT
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                                                                                                                                                                                     Submitted (MAR-1997) to the EMBL/GenBank/DDBJ databases.
-!- SIMILARITY: BELONGS TO THE INTERCRINE ALPHA FAMILY (SMALL CYTOKINE C-x-C) (CHEMOKINE CXC).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
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BY SIMILARITY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 APLATELRCQCLQTLQGIHLKNIQSVKVKSPGPHCAQTEVIATLKNGQKACLNPASPMVK 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi;
Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora; Bovoidea;
Bovidae; Bovinae; Bos.
                                                                                                                     Cetartiodactyla; Ruminantia; Pecora; Bovoidea;
                                                                                                          Chordata; Craniata; Vertebrata; Euteleostomi;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ö
                                               15-DEC-1998 (Rel. 37, Created)
15-DEC-1998 (Rel. 37, Last sequence update)
15-DEC-1998 (Rel. 37, Last annotation update)
GROWTH REGULATED PROTEIN HOMOLOG ALPHA PRECURSOR (GRO-ALPHA).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 104;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         15-DEC-1998 (Rel. 37, Created)
15-DEC-1998 (Rel. 37, Last sequence update)
15-DEC-1998 (Rel. 37, Last annotation update)
GROWTH REGULATED PROTEIN HOMOLOG BETA PRECURSOR (GRO-BETA).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           11; Indels
                                                                                                                                                                                                                                                                                                                                                                           PRINTS; PRO0436; INTERLEDKINB.
PRINTS; PR00437; SMALLCYTKCXC.
PROSITE; PS00471; SMALL_CYTOKINES_CXC; 1.
CYTOKine; Growth factor; Inflammatory response; Signal.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1002CAC064DBlF76 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 299; DB 1;
Pred. No. 2.8e-27;
7; Mismatches 11;
                              104 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     104 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                           SIMILARITY
                                                                                                                                                                                                                                                                                                                                                                                                                          POTENTIAL.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                             66 BY
82 BY
10984 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    78.98;
75.38;
                                                                                                                                                                                                                                                                                                                      EMBL; U95812; AAB93928.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                              SEQUENCE FROM N.A.
Yoshimura T., Modi W.S.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STANDARD;
                              STANDARD;
                                                                                             Bos taurus (Bovine).
Eukaryota; Metazoa; Chor
Mammalia; Eutheria; Ceta
Bovidae; Bovinae; Bos.
NCBI_TaxID=9913;
                                                                                                                                                                                                                                                                                                                                  InterPro; IPR001089; .. InterPro; IPR001811; -.
                                                                                                                                                                                                                                                                                                                                                       InterPro; IPR002473; -. Pfam; PF00048; IL8; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               KIIDKMLNKASAN 104
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           61 KIIEKMLKNGKSN 73
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Bos taurus (Bovine).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   104 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NCBI_TaxID-9913;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Simi
Matches 55;
                             GROA_BOVIN
046676;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GROB_BOVIN
046677;
                                                                                                                                                                                                                                                                                                                                                                                                                                                          DISULFID
                                                                                                                                                                                                                                                                                                                                                                                                                                               DISULFID
                                                                                                                                                                                                                                                                                                                                                                                                                      SIGNAL
                 GROA_BOVIN
                                                                                                                                                                                                                                                                                                                                                                                                                                     CHAIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GROB_BOVIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               92
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Yoshimura T., Modi W.S.;
Submitted (MAR-1997) to the EMBL/GenBank/DDBJ databases.
-!- SIMILARITY: BELONGS TO THE INTERCRINE ALPHA FAMILY (SMALL CYTOKINE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STRAIN=NEW ZEALAND WHITE; TISSUE=Alveolar macrophage;
MEDLINE-95129889; PubMed-7828903;
Johnson M.C., Goodman R.B. II, Kajikawa O., Wong V.A., Mongovin S.M.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 APLATELRCQCLQTLQGIHLKNIQSVKVKSPGPHCAQTEVIATLKNGQKACLNPASPMVK 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GROWTH REGULATED PROTEIN HOMOLOG BETA. BY SIMILARITY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Jose P.J., Collins P.D., Perkins J.A., Beaubien B.C., Totty N.F., Waterfield M.D., Hsuan J., Williams T.J.; Identification of a second neutrophil-chemoattractant cytokine generated during an inflammatory reaction in the rabbit peritoneal cavity in vivo. Purification, partial amino acid sequence and structural relationship to melanoma-growth-stimulatory activity.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     01-JUL-1993 (Rel. 26, Created)
01-FEB-1996 (Rel. 33, Last sequence update)
01-FEB-1998 (Rel. 37, Last annotation update)
PERMEABILITY FACTOR 2 (RFF2) (FRAGMENT).
Oryctolagus cuniculus (Rabbit).
Eukaryota, Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Lagomorpha; Leporidae; Oryctolagus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  .;
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Gene 151:337-338(1994).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 104;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               7; Mismatches 11; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRINTS; PRO0436; INTERLEUKINB.
PRINTS; PRO0437; SMALLCYTKCXC.
PROSTIE; PSO0471; SMALL_CYTOKINES_CXC; 1.
CYTOKINE; Growth factor; Inflammatory response; Signal.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           40A8C06A64D67F7B CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STRAIN-NEW ZEALAND WHITE; TISSUE-Peritoneal cavity; MEDLINE-91378900; PubMed-1898341;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   78.9%; Score 299; DB 1; 75.3%; Pred. No. 2.8e-27;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   71 AA.
                                                                                                                                                                                                                                                                                                                                        or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      POTENTIAL.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        82 B
10950 MW;
                                                                                                                                                                                                                                                                                                                                                                                                       EMBL; U95813; AAB93929.1; -.
                                                                                        C-X-C) (CHEMOKINE CXC).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Best Local Similarity 75.39
Matches 55; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                                                   InterPro; IPR001089; -. InterPro; IPR001811; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               104
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     |||:||| :|
92 KIIDKMLNKASAN 104
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   61 KIIEKMLKNGKSN 73
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               InterPro; IPR002473;
Pfam; PF00048; IL8; 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        104 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE OF 1-36.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NCBI_TaxID=9986;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Martin T.R.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GRO1_RABIT
P30782;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DISULFID
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send an email to license@isb-sib.ch)
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                                                             EMBL; U12310; AAA20487.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE OF 36-88 FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       54; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SCYB1 OR GRO.
Rattus norvegicus (Rat).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STANDARD;
                                                                                                               InterPro; IPR001089; -. InterPro; IPR001811; -.
                                                                                                                                                                             InterPro; IPR002473; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            |||:||| :|
92 KIIDKMLNKASAN 104
                                                                                                                                                                                                    Pfam; PF00048; IL8; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              61 KIIEKMLKNGKSN 73
                                                                                                                                                                                                                                                                                                                                                                                                                                                             104 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE OF 25-96.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NCBI_TaxID=10116;
                                                                                          P09341
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       fibroblasts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TISSUE=Lund
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PROTEIN KC)
                                                                                                                                                                                                                                                                                                                                                                                                                                   DISULFID
SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                         DISULFID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GRO_RAT
P14095;
                                                                                                                                                                                                                                                                                                                                                    SIGNAL
                                                                                                                                                                                                                                                                                                                                                                           CHAIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
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                                                                                                                                                                                                                         This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation-the Buropean Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/
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-!- FUNCTION: PLAYS A ROLE IN MONOCYTE ADHESION TO THE ENDOTHELIUM.
-!- SIMILARITY: BELONGS TO THE INTERCRINE ALPHA FAMILY (SMALL CYTOKINE
                                                                               -!- SUBUNIT: HOMODIMER (PROBABLE).
-!- INDUCTION: GENERATED DURING AN INFLAMMATORY REACTION.
-!- SIMILARITY: BELONGS TO THE INTERCRINE ALPHA FAMILY (SMALL CYTOKINE C-X-C) (CHEMOKINE CXC).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  5 TELRCQCLQTLQGIHLKNIQSVKVKSPGPHCAQTEVIATLKNGQKACLNPASPMVKKIIE 64
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        01-FEB-1996 (Rel. 33, Created)
01-FEB-1996 (Rel. 33, Last sequence update)
01-FEB-1996 (Rel. 33, Last annotation update)
GROWTH REGULATED PROTEIN HOWOLOG PRECURSOR (GRO HOMOLOG).
Oryctolagus cuniculus (Rabbit).
Eukaryota, Metazoa, Chordata; Craniata; Vertebrata; Euteleostomi;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Schwartz D., Chaverri-Alamada L., Berliner J., Kirchgessner T., Quisomoro D., Fang J., Tekamp-Olson P., Lusis J., Fogelman A., Territo M.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ;
0
                             !- FUNCTION: HAS CHEMOTACTIC ACTIVITY FOR NEUTROPHILS (BY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 71;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mammalia; Eutheria; Lagomorpha; Leporidae; Oryctolagus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      E91387CB382C8008 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 296; DB 1; L
Pred. No. 4.1e-27;
9; Mismatches 6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PIR; $1/30/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $1.70/; $
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N -> S (IN REF. 2).
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J. 278:493-497(1991)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match 78.1%;
Best Local Similarity 78.3%;
Matches 54; Conservative 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                          EMBL; L19157; AAA64357.1; -.
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P47854;
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NON_TER
SEQUENCE
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MEDLINE-93246259; PubMed-8482545;
Konishi K., Tanaka Y., Yamamoto M., Yomogida K., Watanabe K.,
Tsurutuji S., Fuljoka M.;
Tsurutuji S., Fuljoka M.;
"Structure of the gene encoding rat neutrophil chemo-attractant Gro.";
Gene 126:285-286(1993).
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Rattus.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            01-NOV-1995 (Rel. 32, Last sequence update)
01-OCT-2000 (Rel. 40, Last annotation update)
GROWTH REGULATED PROTEIN PRECURSOR (CYTOKINE-INDUCED NEUTROPHIL
CHEMOATTRACTANT) (CINC-1) (PLATELET-DERIVED GROWTH FACTOR-INDUCIBLE
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Huang S., Paulauskis J., Kobzik L.;
"Rat KC cDNA cloning and mRNA expression in lung macrophages and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MEDLINE-90062049; PubMed-2684956; Watanabe K., Konishi K., Fujioka M., Kinoshita S., Nakagawa H.; Watanabe K., Konishi K., Fujioka M., Kinoshita S., Nakagawa H.; The neutrophil chemoattractant produced by the rat kidney epithelioid cell line NRK-52E is a protein related to the KC/gro
                                                                                                                                            GROWTH REGULATED PROTEIN HOMOLOG. BY SIMILARITY.
                                                                                                                                                                                                                                                                                                                                    Length 104;
                                                                                                                                                                                                                                                                                                                             Score 295; DB 1; Length 10
Pred. No. 7.9e-27;
8; Mismatches 11; Indels
PRINTS; PRO0436; INTERLEUKINB.
PRINTS; PRO0431; SMALLCYTRCXC.
PROSTIE: PSO0471; SMALL_CYTOKINES_CXC; 1.
Cytokine; Growth factor; Inflammatory response; Signal.
                                                                                                                                                                                                             BY SIMILARITY.
10B9D07B65C77F67 CRC64;
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10900 MW;
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the Buropean Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    --- FUNCTION: HAS CHEMOTACTIC ACTIVITY FOR NEUTROPHILS. CONTRIBUTES TO NEUTROPHIL ACTIVATION DURING INFLAMMATION.
--- SUBUNIT: NONOMER AND HOMODIMER.
--- TISSUE SPECIFICITY: AT LEAST EXPRESSED IN THE LUNG AND TRACHEA.
--- INDUCTION: IN LUNG, BY LIPOPOLYSACCHARIDE OR INFLAMMATION.
--- SIMILARITY: BELONGS TO THE INTERCRINE ALPHA FAMILY (SMALL CYTOKINE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Hanzawa H., Haruyama H., Watanabe K., Tsurufuji S.;
"The three dimensional structure of rat cytokine CINC/Gro in solution
by homonuclear 3D NMR.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Hanzawa H., Haruyama H., Konishi K., Watanabe K., Tsurufuji S.;
"Solution structure of CINC/Gro investigated by heteronuclear NMR.";
J. Biochem. 123:62-70(1998).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Hanzawa H., Haruyama H., Konishi K., Watanabe K., Tsurufuji S.; "Subunit association and monomer structure of CINC/Gro revealed by
Huang S., Paulauskis J.D., Godleski J.J., Kobzik L.;
Expression of macrophage inflammatory protein-2 and KC mRNA in
                                                                                                                                                                                                                                                                                                            a
by
                                                                                                                                                                                                                                                               Nakagawa H., Shiota S., Takano K., Shibata F., Kato H.; "Cytokine-induced neutrophil chemoattractant (CINC)-2 alpha, member of rat GRO/CINCs, is a predominant chemokine produced lipopolysaccharide-stimulated rat macrophages in culture."; Biochem. Biophys. Res. Commun. 220:945-948(1996).
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PRINTS; PRO0437; SMALLCYTKCXC.
PROSITE; PSO0471; SMALL_CYTOKINES_CXC; 1.
Cytokine; Growth factor; Inflammatory response; Signal.
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BY SIMILARITY.

6046C9E2C2582652 CRC64;
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EMBL; M86536; AAA42053.1; --
EMBL; S45856; -; NOT_ANNOTATED_CDS.
PIR; A34481; A34481.
                                                                                                                                                                                                                                 MEDLINE=96183056; PubMed=8607872;
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                                                                    pulmonary inflammation.";
Am. J. Pathol. 141:981-988(1992).
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InterPro; IPR001089; -.
InterPro; IPR001811; -.
InterPro; IPR002473; -.
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                                                                                                                                                                   SEQUENCE OF 25-56.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    J. Immunol. 155:3593-3600(1995).
-!- FUNCTION: HAS CHEMOTACTIC ACTIVITY FOR NEUTROPHILS. CONTRIBUTES TO NEUTROPHIL ACTIVATION DURING INFLAMMATION (BY SIMILARITY).
-!- INDUCTION: BY PLATELET-PERRYDE GROWTH FACTOR. IN LUNG, BY LIPPOPOLYSACCHARIDE OR INFLAMMATION (BY SIMILARITY).
-!- SIMILARITY: BELONGS TO THE INTERCRINE ALPHA FAMILY (SMALL CYTOKINE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Oquendo P., Alberta J., Wen D., Graycar J.L., Derynck R., Stiles C.D., "The platelet-derived growth factor-inducible KC gene encodes a secretory protein related to platelet alpha-granule proteins."; J. Biol. Chem. 264:4133-4137(1989).
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                  01-OCT-1989 (Rel. 12, Last sequence update)
01-OCT-2000 (Rel. 40, Last annotation update)
GROWTH REGULATED PROTEIN PRECURSOR (PLATELET-DERIVED GROWTH FACTOR-INDUCTBLE PROTEIN KC) (SECRETORY PROTEIN N51).
SCYBI OR GRO OR MGSA.
control LPS-induced KC gene transcription in mouse macrophages."; J. Immunol. 155:3593-3600(1995).
                                                                                                                                                                                                                                                                                                                                                                                                 Euteleostomi;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Obmori Y., Fukumoto S., Hamilton T.A., "Two structurally distinct kappa B sequence motifs cooperatively
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Ryseck R.P., Macdonald-Bravo H., Mattel M.-G., Bravo R.; "Cloning and sequence of a secretory protein induced by growth
                                                                                                                                                                                                                                                                                                                                                                                             Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleosto
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             or send an email to license@isb-sib.ch).
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EMBL; U20634; AAB03376.1; -.
EMBL; U20527; AAB03376.1; JOINED.
EMBL; S79767; -; NOT_ANNOTATED_CDS.
PIR; A32954; A32954.
PIR; JH0081; JH0081.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         factors in mouse fibroblasts.";
Exp. Cell Res. 180:266-275(1989).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MEDLINE=96016008; PubMed=7561058;
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                                                                                                                                                                                                                    STANDARD;
                                                                                                                                                                                                                                                                                                                                                                               Mus musculus (Mouse)
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85 KIVQKMLK 92
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P12850:
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51; Conservative

Query Match Best Local Similarity Matches 51; Conserv

74.98; 75.0%;

Length 96;

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EMBL; U95809; AAB93925.1; -.
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055235;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             -!- SUBUNIT: HOMOTETRAMER.
-!- SIMILARITY: BELONGS TO THE INTERCRINE ALPHA FAMILY (SMALL CYTOKINE
                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 APLATELRCQCLQTLQGIHLKNIQSVKVKSPGPHCAQTEVIATLKNGQKACLNPASPMVK 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Wolpe S.D., Sherry B., Juers D., Davatelis G., Yurt R.W., Cerami A.; "Identification and characterization of macrophage inflammatory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Shao W., Jerva L.F., West J., Lolis E., Schweitzer B.I.; "Sollution structure of murine macrophage inflammatory protein-2."; Blochemistry 37:8303-8131(1998).
-i. FUNCTION: CHEMOTACTIC FOR HUMAN POLYMORPHONUCLEAR LEUKOCYTES BUT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mus musculus (Mouse).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
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Tekamp-Olson P., Gallegos C., Bauer D., McClain J., Sherry B. Fabre M., van Deventer S., Cerami A.;
"Cloning and characterization of cDNAs for murine macrophage inflammatory protein 2 and its human homologues.";
J. Exp. Med. 172:911-919(1990).
                                                                                                                                                                                                                                                                                                                                                                        73.4%; Score 278; DB 1; Length 96; 73.5%; Pred. No. 6.2e-25; ive ll; Mismatches 7; Indels
                             MAUJ. MALILLONGOS, GIOLI.
INTERPRO; IPR001089; -
INTERPRO; IPR001811; -
INTERPRO; IPR002473; -
Pfam; PF00048; IL6; 1.
PRINTS; PR00436; INTERLEUKINB.
PRINTS; PR00471; SMALLCYTKCXC.
PROSITE; PS00471; SMALLCYTKCXC.
CYTOKINE; Growth factor; Inflammatory response; Signal.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DOES NOT INDUCE CHEMOKINESIS OR AN OXIDATIVE BURST
                                                                                                                                                                                                                                  GROWTH REGULATED PROTEIN.
BY SIMILARITY.
BY SIMILARITY.
                                                                                                                                                                                                                                                                                                       4A52B5E5C38B45C2 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        01-JUL-1989 (Rel. 11, Created)
01-FEB-1991 (Rel. 17, Last sequence update)
30-MAY-2000 (Rel. 39, Last annotation update)
MACROPHAGE INFLANMATORY PROTEIN 2 PRECURSOR (MIP2).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Proc. Natl. Acad. Sci. U.S.A. 86:612-616(1989)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      100 AA.
                                                                                                                                                                                                                    PROBABLE
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                                                                                                                                                                                                                                                                                                       96 AA; 10254 MW;
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Best Local Similarity 73.5%,
Matches 50; Conservative
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                      MGD; MGI:108068; Grol.
P09341; 1MSH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE FROM N.A.
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P10889;
                                                                                                                                                                                                                                                             DISULFID
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use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
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-!- FUNCTION: HAS CHEMOTACTIC ACTIVITY FOR NEUTROPHILS.
-!- SIMILARITY: BELOWGS TO THE INTERCRINE ALPHA FAMILY (SMALL CYTOKINE C-X-C) (CHEMOKINE CXC).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               28 AVVASELRCQCLKTLPRVDFKNIQSLSVTPPGPHCAQTEVIATLKGGGKVCLDPEAPLVQ 87
                                                                                                                                                                                                                                                                                                                                                                      InterPro; IPR002473; --
Pfam; PF00048; IL8; 1.
PRINTS; PR00436; INTERLEUXINB.
PRINTS; PR00436; SMALLCYRCXC.
PROSITE; PS00471; SMALL_CYTOKINBS_CXC; 1.
CYtokine; Chemotaxis; Inflammatory response; Signal; 3D-structure.
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Mammalia; Eutheria; Rodentia; Hystricognathi; Caviidae; Cavia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MACROPHAGE INFLAMMATORY PROTEIN 2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 68.9%; Score 261; DB 1; Length 100; 65.8%; Pred. No. 5.5e-23;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     13; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     15-DEC-1998 (Rel. 37, Created)
15-DEC-1998 (Rel. 37, Last sequence update)
10-CT-2000 (Rel. 40, Last annotation update)
GROWTH REGULATED PROTEIN PRECURSOR.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         104 AA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SCYBl OR GRO. Cavia porcellus (Guinea pig).
                                                                                                                                                                   EMBL; X53798; CAA37807.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            48; Conservative
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Yoshimura T., Modi W.S.;
                                                                                                                                                                                                                                      PDB; IMI2; 29-APR-98.
MGD; MGI:1340094; Scyb2.
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                                                                                                                                                                                                                                                                                                         InterPro; IPR001089; -. InterPro; IPR001811; -.
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MEDLINE-94318061; PubMed-8043001;
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                                                                                                                                                                                                                                                                                                                                                                             Rattus norvegicus (Rat).
Bukaryota: Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Rattus.
NCBL_TaxDD=10116;
                                                                                                                                                                                         Gaps
                                                                                                                                                                                                         1 APLATELRCQCLQTLQGIHLKNIQSVKVKSPGPHCAQTEVIATLKNGQKACLNPASPMVK 60
                                                                                                                                                                                                                   Feng L., Xia Y., Yoshimura T., Wilson C.B.;
Modulation of neutrophil influx in glomerulonephritis in the rat
with anti-macrophage inflammatory protein-2 (MIP-2) antibody.";
J. Clin. Invest. 95:1009-1017(1995).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Huang S., Paulauskis J.D., Godleski J.J., Kobzik L.; "Expression of macrophage inflammatory protein-2 and RC mRNA in
                                                                                                                                                                                         ö
                                                                                                                                                                                                                                                                                                                               01-APR-1993 (Rel. 25, Created)
01-APR-1993 (Rel. 25, Last sequence update)
01-OCT-2000 (Rel. 40, Last annotation update)
MACROPHAGE INFLAMATORY PROTEIN 2 PRECURSOR (MIP2) (CINC-3).
SCYB2 OR MIP2 OR MIP-2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE FROM N.A.
STRAIN=CD; TISSUE=Lung;
Farone A., Farone M., Shi M.M., Kobzik L., Paulauskis J.D.;
Submitted (MAR-1996) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                    67.8%; Score 257; DB 1; Length 104; 61.1%; Pred. No. 1.6e-22; 1ive 19; Mismatches 9; Indels
                                 Submitted (APR-1992) to the EMBL/GenBank/DDBJ databases.
                                                                                                                      BY SIMILARITY.
BY SIMILARITY.
DAFD96D9F4041D5A CRC64;
                                                                                                             GROWTH REGULATED PROTEIN
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                                                                                                       POTENTIAL.
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MEDLINE=95189993; Pubmed=7883948;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              monary inflammation.";
J. Pathol. 141:981-988(1992).
                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE FROM N.A.
STRAIN-FISCHER 344; TISSUE-Lung;
EMBL; U95810; AAB93926.1; -.
EMBL; AF052004; AAC06227.1; -.
EMBL; AF052005; AAC06228.1; -.
InterPro; IPR001089; -.
                                                                                                                                          11069 MW;
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                                                                                                                                                                                        44; Conservative
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66
82
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KVLQRMLKGSKA 103
                                                                                                                                                                                                                                              61 KIIEKMLKNGKS 72
                                                                                                                                         104 AA;
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SEQUENCE OF 32-100.
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Matches 44; Conserv
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DISULFID
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                                                                                                                                                                                                                                                                   Nakagawa H., Komorita N., Shibata F., Ikesue A., Konishi K., Fujioka M., Kato H.;
"Identification of cytokine-induced neutrophil chemoattractants
(CINC), rat GRO/CINC-2 alpha and CINC-2 beta, produced by granulation tissue in culture: purification, complete amino acid sequences and characterization."
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         30 VASELRCQCLTTLPRVDFKNIQSLTVTPPGPHCAQTEVIATLKDGHEVCLNPEAPLVQRI 89
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7
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PRINTS: PR00437; SMALLCYTKXKLE.
PROSTTE: P800471; SMALLCYTKXKLE.
CYTCKINE; Chemotaxis; Inflammatory response; Signal.
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EMBL; U45965; AAA92438.1; -...
EMBL; S45855; -; NOT_ANNOTATED_CDS.
PIR; S21467; S21467.
HSSP; P10889; 1MI2.
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78 B
10783 MW;
                                                                                                                                                 Biochem. J. 301:545-550(1994).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       EMBL; X65647; CAA46599.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               C-X-C) (CHEMOKINE CXC).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity 62.09
Matches 44; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               InterPro; IPR001089; -. InterPro; IPR001811; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 InterPro; IPR002473; -. Pfam; PF00048; IL8; 1.
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38
100 AA;
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                                                                                                                                                                                                SEQUENCE OF 32-59.
                                                                                                                             characterization.
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us-09-467-160-3.rsp

GenCore version 4.5 Copyright (c) 1993 - 2000 Compugen Ltd.

OM protein - protein search, using sw model

May 4, 2001, 07:35:28; Search time 19.46 Seconds (without alignments) 439.680 Million cell updates/sec Run on:

US-09-467-160-3
379
1 APLATELRCQCLQTLQGIHL.....PASPMVKKIIEKMLKNGKSN 73 Title: Perfect score: Sequence:

BLOSUM62 Gapop 10.0 , Gapext 0.5 Scoring table:

374700 seqs, 117207915 residues hrched:

374700 Total number of hits satisfying chosen parameters:

Minimum DB seq length: 0 Maximum DB seq length: 200000000

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

SPTREMBL_15:* Database :

1: sp_archea:*
2: sp_barchea:*
3: sp_lungi:*
4: sp_human:*
5: sp_invertebrate:*
6: sp_mammal:*
7: sp_mho:*
8: sp_organelle:*
9: sp_lange:*
10: sp_lange:*
11: sp_ordangi:*
12: sp_unclassified:*
13: sp_vertebrate:*
14: sp_vertebrate:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

STIMMARTES

	Description	Q28724 oryctolagus	062764 equus cabal	073912 gallus gall	Q9z126 mus musculu	Q9xsx5 felis silve	Q9quq7 rattus sp.	Q9uc64 homo sapien	Q9tr91 bos taurus	Q9yva9 qa11id herp	Q9ibj7 turkey herp	Q67634 qallid herp	097800 bos taurus	093442 lampetra f1	Q9qv88 rattus sp.	Q9qv74 rattus sp.	09xsd2 oryctolagus	Q9qx28 mus musculu	Q9jhh7 mus musculu	Q9ns21 homo sapien
SUMMARIES	ID	028724	062764	073912	Q9Z126	09xsx5	090007	Q9UC64	Q9TR91	Q9YVA9	Q91BJ7	067634	097800	093442	090V88	090V74	09xsD2	090x28	09лнн7	Q9NS21
	DB	9	9	13	11	9	11	4	9	14	14	14	9	13	11	11	9	11	11	4
	% Query Match Length DB	108	59	104	105	101	33	54	48	134	142	203	31	101	34	40	41	86	66	111
	% Query Match	79.2	63.6	47.4	45.0	37.3	36.1	35.1	34.0	33.1	33.1	33.1	32.5	30.9	28.5	28.2	27.8	23.9	23.4	23.1
	Score	300	241	179.5	170.5	141.5	137	133	129	125.5	125.5	125.5	123	117	108	107	105.5	90.5	88.5	87.5
	Result No.	1	7	М	4	2	9	7	æ	6	10	11	12	13	14	15	16	17	18	19

09jhh5 mus musculu 09qzdl rattus norv 068406 human cytom 09qv82 rattus sp. 09j2ml macaca mula 06339 human cytom 023352 caenorhabdi 09ptf8 brachydanio 09quib rattus sp. 09quib rattus sp. 09quib rattus sp. 09quib rattus sp.	090825 gallus gall 090825 gallus gall 09wrt7 macaca mula 09u5yl dictyostell 082651 jembrana di 09qy73 rattus sp. 09u73 rattus sp. 09u73 matus sp.	Q98158 kaposi's sa O6045 homo sapien O81815 arabidopsis O96437 elmeria ten Q91426 box taurus Q9445 box taurus Q29443 archaeoglob Q9x6x2 myxococcus O92368 vesv-like c
11 09JHH5 11 090ZD1 11 090ZD1 11 090V82 14 068399 14 068399 15 023352 13 090FTF8		0 0 000
		95 1 471 4 397 1 617 5 1212 1 397 6 349 1 416 2
200.6 19.7 19.5 18.9 18.3 18.3	18.1 17.9 17.2 17.0 16.6 16.4	16.00
78 74.5 74.5 71.5 71.5 71.5 69.5	68 69 60 60 60 60 60 60 60 60	61.5 61.5 61 60.5 60.5 60.5
22 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2	9 9 9 9 9 9 9 9 9 9 9 9 9 9 9 9 9 9 9	2 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8

ALIGNMENTS

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Sick C., Schneider K., Staehell P., Weining K.C.; "Novel chicken CXC and CC chemokines."; Cytokine 12:181-186(2000).
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Matches 32; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PFAM; PF00048; IL8; 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Best Local Similarity
Matches 36; Conserve
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  63 IEKMLKNGKSN 73
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 89 IKAILDKADTN 99
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NCBI_TaxID=10090;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
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Q92126;
                                                                                                                                                                                                                                                                                                                                                                                                                                          Signal
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Q92126
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Archosauria; Aves; Neognathae; Galliformes; Phasianidae; Phasianinae;
                                                                          Gaps
                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      15 LQGIHLKNIQSVKVKSPGPHCAQTEVIATLKNGQKACLNPASPMVKKIJEKMLKNGKSN 73
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Eŭkaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Perissodactyla, Equidae, Equus.
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                       Length 108;
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                                                                          Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TISSUE=SKIN;
Rieder S., Checa-Cortes M.L., Joerg H., Stranzinger G.;
Submitted (MAR.1998) to the EMBL/GenBank/DDBJ databases.
HMBL; AF053497; AAC97075.1;
HSSP; 1918975; 10NK
INTERPRO; IPR001089; -.
INTERPRO; IPR001811; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Last sequence update)
Last annotation update)
ACTIVITY HOMOLOG (FRAGMENT).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        59 AA; 6344 MW; 1CF4CF77549C0CE4 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               01-AUG-1998 (TrEMBLrel. 07, Created)
01-AUG-1998 (TrEMBLrel. 07, Last sequence update)
01-CCT-2000 (TrEMBLrel. 15, Last annotation update)
K60 PROTEIN PRECURSOR (CXC CHEMOKINE K60).
Score 300; DB 6; Le
Pred. No. 1.4e-27;
.......
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78.0%; Pred. No. 5.5e-21;
Live 5; Mismatches 8;
                                                                                                                                                                                                                                                                                                                                                                                                            59 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     104 AA
                                                                    8; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Created)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PFAM, PF00048; ILB; 1.
PRINTS; PR00436; INTERLEUKINB.
PRINTS; PR00437; SMALLCYTKCXC.
                  ch 79.2%;
1 Similarity 76.7%;
56; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                            01-AUG-1998 (TrEMBLrel. 07,
01-AUG-1998 (TREMBLrel. 07,
01-OCT-2000 (TREMBLRel. 15,
MELANOMA GROWTH STIMULATORY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match 63.6
Best Local Similarity 78.0
Matches 46; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                               PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRELIMINARY;
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                                                                                                                                                                                                                                                        | ::| | | ||
96 KFLQKRLSKGSSN 108
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Equus caballus (Horse)
                                                                                                                                                                                                                           61 KIIEKMLKNGKSN 73
                     Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE FROM N.A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NCBI_TaxID=9796;
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SEQUENCE
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                                                                          Matches
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OOC SCHOOL SCHOO

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Watanabe O., Tamari M., Natori K., Kubo S., Shiomoto Y., Nakamura Y.;
"Isolation of the murine homologue of rat and human PF4platelet factor AcDNA from the spleen of NOA mouse that is a new atopic dermatitis model by a differential display method.";
Submitted (SEP-1998) to the EMBL/Genbank/DDBJ databases.
EMBL; AB017491; BAA75660.1; -.
HSSP; P02776; IRHP.
MGD; MGI:1889711; Pf4.
INTERPRO; IPR001089; -.
INTERPRO; IPR001811; -.
INTERPRO; IPR001473; -.
BEQUENCE FROM N.A.

Hughes S.M., Bunstead N.;

Hughes S.M., Bunstead N.;

Mapping of a second ELR CXC chemokine to chicken chromosome four.";

Submitted (JUN 2000) to the EMBL/GenBank/DDBJ databases.

EMBL; Y14971; CAA75212.1; -.

EMBL; AR277660; AAF86485.1; -.

HSSP; P02775; ITVX.

INTERPO; IPR00189; -.

PRNM: PR0048; IL8; 1.

PRINTS; PR00437; SMALLCYTKCXC.

PROSITE; PS00471; SMALL_CYTOKINES_CXC; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    4 ATELRCQCLQT-LQGIHLKNIQSVKVKSPGPHCAQTEVIATLKNGQKACLNPASPMVKKI 62
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Ĥ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 104;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        47.4%; Score 179.5; DB 13; Length ilarity 50.7%; Pred. No. 1.4e-13; Conservative 13; Mismatches 21; Indels
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                                                                                                                                                                                                                                                                                                                                               POTENTIAL.
K60 PROTEIN.
83725AB91FE18E3D CRC64;
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SEQUENCE 105 AA; 11243 MW; 94F0190EFACF1A6C CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Last sequence update)
Last annotation update)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       14;
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49.2%; Pred. No. 1.6e-12;
Live 18; Mismatches 14,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         105 AA
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21 104 K
104 AA; 11199 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE FROM N.A.
STRAIN=C57BL6J; TISSUE=SPLEEN;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRINTS; PRO0436; INTERLEUKINB
PRINTS; PRO0437; SMALLCYTKCXC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      01-MAY-1999 (TrEMBLrel. 10, 01-MAY-1999 (TrEMBLrel. 10, 01-OCT-2000 (TrEMBLrel. 15, PLATELET FACTOR 4.
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Fri May

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INTERPRO; IPR001089; -.
INTERPRO; IPR001811; -.
INTERPRO; IPR002473; -.
PFAM; PF00048; IL8; 1.
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                                                                                                             Query Match
Best Local Similarity
Matches 25; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 54 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NCBI_TaxID=9913;
                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE
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                                                                                                                                                                                                                                                  Q9UC64
Q9UC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Q9TR91;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Q9TR91
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Nakagawa H., Ikesue A., Hatakeyama S., Kato H., Gotoda T.,
Komorita N., Watanabe K., Miyai H.;
Production of an interleukin-8-like chemokine by cytokine-stimulated
rat NRK-49F fibroblasts and its suppression by anti-inflammatory
              Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Rattus.
NCBL_TaxID=10118;
6 ELRCQCLQTL-QGIHLKNIQSVKVKSPGPHCAQTEVIATLKNGQKACLNPASPMVKKIIE 64
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           28 ISSELRCQCIKTHSTPFNPKLIKELTVIDSGPHCENSEIIVKLVNGKEVCLDPKQKWVQK 87
                                                                                                                                                                                                            Felis silvestris catus (Cat).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Carnivora; Fissipedia; Felidae; Felis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              6; Length 101;
                                                                                                                                                                                                                                                                                   Straubinger A.F., Simpson K.W., Straubinger R.K.;

Straubinger A.F., Simpson K.W., Straubinger R.K.;

Submitted (JUN 1999) to the EMBL/GenBank/DDBJ databases.

EMBL; AF158598; AAD40323.1; -.

REMED; AF158599; AAD40323.1; -.

INTERPRO; IPR001089; -.

INTERPRO; IPR00189; -.

INTERPRO; IPR00143; -.

PEAM; PF00048; IL8; 1.

RINTERPO; IPR00436; IL8; 1.

REMINTS; PR00436; IL8; 1.

REMINTS; PR00437; SMALLCYTKCXC.

REMINTS; PR00437; SMALLCYTKCXC.

REMINTS; PR00437; SMALLCYTYCXINES_CXC; 1.

SEQUENCE 101 AA; 11165 MW; 690D97F13EF79170 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
                                                                                                                                                          01-NOV-1999 (TrEMBLrel. 12, Created)
01-NOV-1999 (TrEMBLrel. 12, Last sequence update)
01-OCT-2000 (TrEMBLrel. 15, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        01-MAY-2000 (TrEMBLrel. 13, Created)
01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-OCT-2000 (TrEMBLrel. 15, Last annotation update)
CHEMOATTRACTANT P-3 (FRAGMENT).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          37.3%; Score 141.5; DB 6;
38.6%; Pred. No. 3.5e-09;
tive 17; Mismatches 25;
                                                                                                                                     101 AA
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HSSP; P19875; 1QNK.
                                                                                                                                      PRT;
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                                                                                                                                      PRELIMINARY;
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Best Local Similarity
Matches 27; Conserv
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VVEIFLKKAE 97
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                                                                                                                                                                                                                                                  NCBI_TaxID=9685;
                                                                     101 KILES 105
                                                                                                                                                                                                INTERLEUKIN-8.
                                               65 KMLKN 69
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01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-MAY-2000 (TrEMBLrel. 15, Last annotation update)
01-OCT-2000 (TrEMBLrel. 15, Last annotation update)
PF 4-DERIVED ENDOTHELIAL CELL GROWTH INHIBITOR PEAK II (FRAGMENT).
Homo sapiens (Human).
Eukaryota: Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia: Eutheria; Primates; Catarrhini; Hominidae; Homo.
NCBL_TAXID=9606;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Allmann-Iselin I., Car B.D., Zwahlen R.D., Mueller-Schupbach R., Wader-Walther M., Steckholzer U., Walz A.; Bovine ENR, a new monoscophage derived cytokine of the interleukin-8 family. Structure, function, and expression in acute
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi;
Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora; Bovoidea;
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01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-0CT-2000 (TrEMBLrel. 15, Last annotation update)
EPITHELIAL-DERIVED NEUTROPHIL-ACTIVATING PEPTIDE (FRAGMENT).
                                                                                                                                        Length 33;
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                                                                                                                                                                                            Indels
PRINTS; PRO0436; INTERLEUKINB.
PRINTS; PRO0437; SMALLCYTKCXC.
SEQUENCE 33 AA; 3561 MW; 2740F8ED16C8C1FF CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               C0B560236BF1B14A CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gupta S.K., Hassel T., Singh J.P.;
Proc. Natl. Acad. Sci. U.S.A. 92:7799-7803(1995).
HSSP; P02776; 1RHP.
                                                                                                                                     Score 137; DB 11;
Pred. No. 3.7e-09;
3; Mismatches 5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match 35.1%; Score 133; DB 4; Best Local Similarity 49.0%; Pred. No. 1.8e-08; Matches 24; Conservative 14; Mismatches 11.
                                                                                                                                                                                                                                                                                                                                                                                                                                           54 AA.
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                                                                                                                                                                                                                                            PRT;
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                                                                                                                                        36.1%;
75.8%;
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"Isolation and characterization of Marek's disease virus (MDV) cDNAs mapping to the BamHI-12, BamHI-02, and BamHI-L fragments of the MDV genome from lymphoblastoid cells transformed and persistently infected
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STRAIN=GA;
BODLINE=96074534; PubMed=7491783;
Peng Q., Zeng M., Bhuiyan Z.A., Ubukata E., Tanaka A., Nonoyama M.,
Shirazi Y.;
                                                         MEDLINE-92237304; PubMed=1315048;
Jones D., Lee L., Liu J.L., Kung H.J., Tillotson J.K.;
"Marek disease virus encodes a basic-leucine zipper gene resembli
the fos/jun oncogenes that is highly expressed in lymphoblastoid
                                                                                                                                                                                                                     STRAIN=GA;
Lee L.F., Wu P., Sui D., Ren D., Kung H.J.;
Lee L.F., Wu P., Sui D., Ren D., Kung H.J.;
Late Complete UL Sequence of Serotype I Marek's Disease Virus.";
Proc. Natl. Acad. Sci. U.S.A. 0:0-0(2000).
Brnb1, Arl47806; AAF67205.1;
BRNB1, AF147806; AAF6739.1;
SEQUENCE 142 AA; 15770 MW; 35A8CEB9B29FD209 CRC64;
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33.1%; Score 125.5; DB 14; Length 203;
Best Local Similarity 40.0%; Pred. No. 5.2e-07;
Matches 26; Conservative 16; Mismatches 22; Indels 1;
                                                                                                                                                                                                                                                                                                                                                                                              DB 14; Length 142;
                                                                                                                                                                                                                                                                                                                                                                                          Ouery Match 33.1%; Score 125.5; DB 14; Length Best Local Similarity 40.0%; Pred. No. 3.6e-07; Matches 26; Conservative 16; Mismatches 22; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   01-NOV-1996 (TrEMBLrel. 01, Created)
01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
01-OCT-2000 (TrEMBLrel. 15, Last annotation update)
ECO & PROTEIN (FRAGMENT).
gallid herpesvirus 1.
Viruses; dsDNA viruses, no RNA stage; Herpesviridae;
Alphaherpesvirinae; Varicellovirus.
                                                                                                                                                             Proc. Natl. Acad. Sci. U.S.A. 89:4042-4046(1992)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     203 AA
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VICOLOGY 213:590-599(1995).
EMBL; U34966; AAC54629:1; -.
HSSP; P10889; IMIZ.
INTERPRO; IPR00169; -.
INTERPRO; IPR001811; -.
INTERPRO; IPR001811; -.
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SEQUENCE FROM N.A.
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88 FIKKL 92
                                               STRAIN=GA;
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SEQUENCE
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Q67634
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                                                                                                                                                                                                 Length 48;
                                                                                                                                                                                                                                       Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STRAIN=GA;
Lin S.-F., Robinson D., Chen H.-C., Kung H.-J.;
LCXC chemokine encoded by Marek's Disease Virus (MDV).";
Submitted (MAY-1998) to the EMBL/GenBank/DDBJ databases.
EMBL; AF065430; AAC77449.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRINTS; PRO0436; INTERLEUKINB.
PRINTS; PRO0437; SMALLCYTKCXC.
SEQUENCE 134 AA; 14828 MW; C859CDB868D25190 CRC64;
pulmonary inflammation.";
Am. J. Pathol. 145:1382-1389(1994).
HSSP; PO2775; INAP.
INTERPRO; IPR001089; -.
PRAM; PR00048; IL8; 1.
PRINTS; PR00437; SMALLCYTKCXC.
SEQUENCE 48 AA; 5066 MW; B80EEI5AC522FEA8 CRC64;
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gallid herpesvirus 1.
Viruses; dsDNA viruses, no RNA stage; Herpesviridae;
Alphaherpesvirinae; Varicellovirus.
NCBI_TaxID=10386;
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Viruses; dsDNA viruses, no RNA stage; Herpesviridae.
NCBI_TaxID=10390;
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Last annotation update)
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                                                                                                                                                                                                                                       12;
                                                                                                                                                                                                                                                                                                   Ouery Match 34.0%; Score 129; DB 6; Best Local Similarity 57.1%; Pred. No. 4.7e-08; Matches 24; Conservative 6; Mismatches 12
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R-LORF2.
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SIGNAL
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Q9QV88;
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Q9QV88
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Najakshin A.M., Mechetina L.V., Alabyev B.Y., Taranin A.V.; "Identification of the interleukin 8 homologue in lamprey (Lampetra fluviatilis): early evolutionary divergence of chemokines."; Submitted (JUL.1998) to the EMBL/GenBank/DDBJ databases.
                                                                                                    01-MAY-1999 (TrEMBLrel. 10, Created)
01-MAY-1999 (TrEMBLrel. 10, Last sequence update)
01-OCT-2000 (TrEMBLrel. 15, Last annotation update)
01-OCT-2000 (TREMBLREL. 18, Last annotation update)
EPITHELIAL CELL INFLAMMATORY PROTEIN-1 (ECIP-1) (FRAGMENT).
Bos taurus (Bovine).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora; Bovoidea;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Lampetra fluviatilis (River lamprey).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Hyperoartia;
Petromyzontiformes; Petromyzontidae; Lampetra.
                                                                                                                                                                                                                                                                                                                                                                                                                                             ö
                                                                                                                                                                                                                                                                                                                                                                                                                         32.5%; Score 123; DB 6; Length 31; 74.2%; Pred. No. 1.5e-07; tive 3; Mismatches 5; Indels
                                                                                                                                                                                                                                                                                                                                                                                              31 AA; 3386 MW; 74172DC13D3A72B8 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 01-NOV-1998 (TrEMBLrel. 08, Created)
01-NOV-1998 (TrEMBLrel. 08, Last sequence update)
01-OCT-2000 (TrEMBLrel. 15, Last annotation update)
LFCA-1 PROTEIN PRECURSOR.
                                                                                   31 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 101 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         43 TLKNGQKACLNPASPMVKKIIEKMLKNGKSN 73
                                                                                   PRT;
                                                                                                                                                                                                   SEQUENCE FROM N.A., AND INDUCTION.
TISSUE=KIDNEY EPITHELIUM;
                                                                                                                                                                                                                                                                                                                                            INTERPRO: IPRO01811; -. PFAM: PF00048; ILB; 1. Cytokine; Inflammatory response.
                                                                                                                                                                                                                                                                                                                                                                                                                                             23; Conservative
                                                                                  PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRELIMINARY;
                                                                                                                                                                       Bovidae; Bovinae; Bos.
                                                                                                                                                                                                                                                                                                                                                                                                                                  Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TISSUE=LEUKOCYTES;
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157 FIKKL 161
                  62 IIEKM 66
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SEQUENCE
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Matches 2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDINE-93192044; PubMed-8383510; Driscoll K.E., Hassenbein D.G., Carter J., Poynter J., Asquith T.N., Grant R.A., Whitten J., Purdon M.P., Takigiku R.; "Macrophage inflammatory proteins I and 2: expression by rat alveolar macrophages, fibroblasts, and epithelial cells and in rat lung after mineral dust exposure.";
                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Rattus sp.
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Rattus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                    30 RCQCVHVISKFIHPKHFQTMEVIPQSSNCKNVEIIVTMKSTNNQICLNPDAPWVRKVISH 89
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Rattus.
                                                                                                                                                                                                                                                                                                                                                                                                                     8 RCQCLQTL-QGIHLKNIQSVKVKSPGPHCAQTEVIATLKN-GQKACLNPASPMVKKIIEK 65
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                               5
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                                                                                                                                                                                                                                                                                                      Length 101;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          28.5%; Score 108; DB 11; Length 34; 61.8%; Pred. No. 9e-06;
                                                                                                                                                                                                                                                                                                                                                               Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
                                                                                                                                                                    22 POTENTIAL.
101 LFCA-1 PROTEIN.
11095 MW; 80CFEE81EA7336D2 CRC64;
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01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-OCT-2000 (TrEMBLrel. 15, Last annotation update)
MACROPHAGE INFLAMMATORY PROTEIN 2 HOMOLOG (FRAGMENT).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          34 AA; 3762 MW; ECF6DD4262F44782 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         01-MAY-2000 (TrEMBLrel. 13, Created)
01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-OCT-2000 (TrEMBLrel. 15, Last annotation update)
MACROPHAGE INFLAMMATORY PROTEIN 2 (FRAGMENT).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     6
                                                                                                                                                                                                                                                                                                      Query Match 30.9%; Score 117; DB 13; Best Local Similarity 34.3%; Pred. No. 2.5e-06; Matches 23; Conservative 18; Mismatches 24;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Am. J. Respir. Cell Mol. Biol. 8:311-318(1993).
HSSP; P10889; IMI2.
INTERPRO; IPRO01089; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            40 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     4; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRT;
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HSSP; P02775; 1TVX.
INTERPRO; 1PR001089; -.
INTERPRO; 1PR00181; -.
PFAM; PF00048; ILS; 1.
PRINTS; PR00437; SMALLCYTKCXC.
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PRINTS; PR00437; SMALLCYTKCXC.
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Best Local Similarity 61.8
Matches 21; Conservative
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NCBI_TaxID=10118;

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                               SEQUENCE.
MEDLINE-93285151; PubMed-8508797;
Watanabe K., Iida M., Takaishi K., Suzuki T., Hamada Y., Iizuka Y.,
Tsurufuji S.; Iida M., Takaishi K., Suzuki T., Hamada Y., Iizuka Y.,
Tsurufuji S.;
"Chemoattractants for neutrophils in lipopolysaccharide-induced
inflammatory exudate from rats are not interleukin-8 counterparts but
goro-peroduct/melanoma-growth-stimulating-activity-related
factors.";
Eur. J. Biochem. 214:267-270(1993).
HSSP; P10889; LMI2.
SEQUENCE 40 AA; 4500 MW; 330DFI3BAE51ACFD CRC64;
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